

Network Systems  
Science & Advanced  
Computing  
Biocomplexity Institute  
& Initiative  
University of Virginia

# Estimation of COVID-19 Impact in Virginia

December 22<sup>nd</sup>, 2021

(data current to December 10<sup>th</sup> – 14<sup>th</sup>)

Biocomplexity Institute Technical report: TR 2021-129



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**BIOCOMPLEXITY** INSTITUTE

[biocomplexity.virginia.edu](https://biocomplexity.virginia.edu)

# About Us

- Biocomplexity Institute at the University of Virginia
  - Using big data and simulations to understand massively interactive systems and solve societal problems
- Over 20 years of crafting and analyzing infectious disease models
  - Pandemic response for Influenza, Ebola, Zika, and others



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# Overview

- **Goal:** Understand impact of COVID-19 mitigations in Virginia
- **Approach:**
  - Calibrate explanatory mechanistic model to observed cases
  - Project based on scenarios for next 4 months
  - Consider a range of possible mitigation effects in "what-if" scenarios
- **Outcomes:**
  - Ill, Confirmed, Hospitalized, ICU, Ventilated, Death
  - Geographic spread over time, case counts, healthcare burdens

# Key Takeaways

Projecting future cases precisely is impossible and unnecessary.

Even without perfect projections, we can confidently draw conclusions:

- **Case rates grew after holiday break but growth has slowed slightly, keeping case rates high as the anticipated arrival of Omicron may fuel more rapid growth in the near term**
- VA 7-day mean daily case rate up to 38.5/100K from 30/100K; US is up to 44/100K (from 36/100K)
- Projections show a continued rise of cases which becomes more extreme under Omicron and FallWinter scenarios that anticipate likely drivers of future transmission
- Recent updates:
  - Overhauled model structure further refined to better capture different tiers of immunity and the immune evasion of the Omicron variant
  - Analysis of the effects of increasing 3<sup>rd</sup> dose coverage

The situation continues to change. Models continue to be updated regularly.



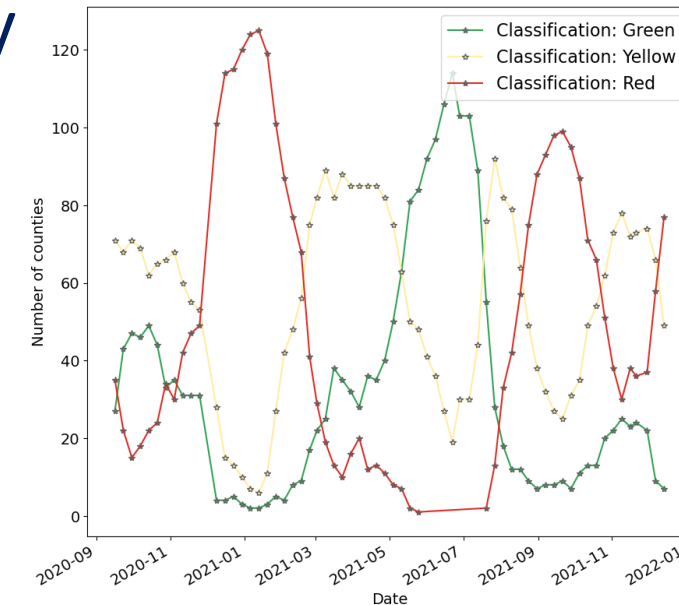
# Situation Assessment

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# Case Rates (per 100k) and Test Positivity

- Case rate increase across all health districts
- Some past 50% of winter peak and growing
- More than 50% of counties with TPR > 10%

Data source: <https://data.cms.gov/covid-19/covid-19-nursing-home-data>

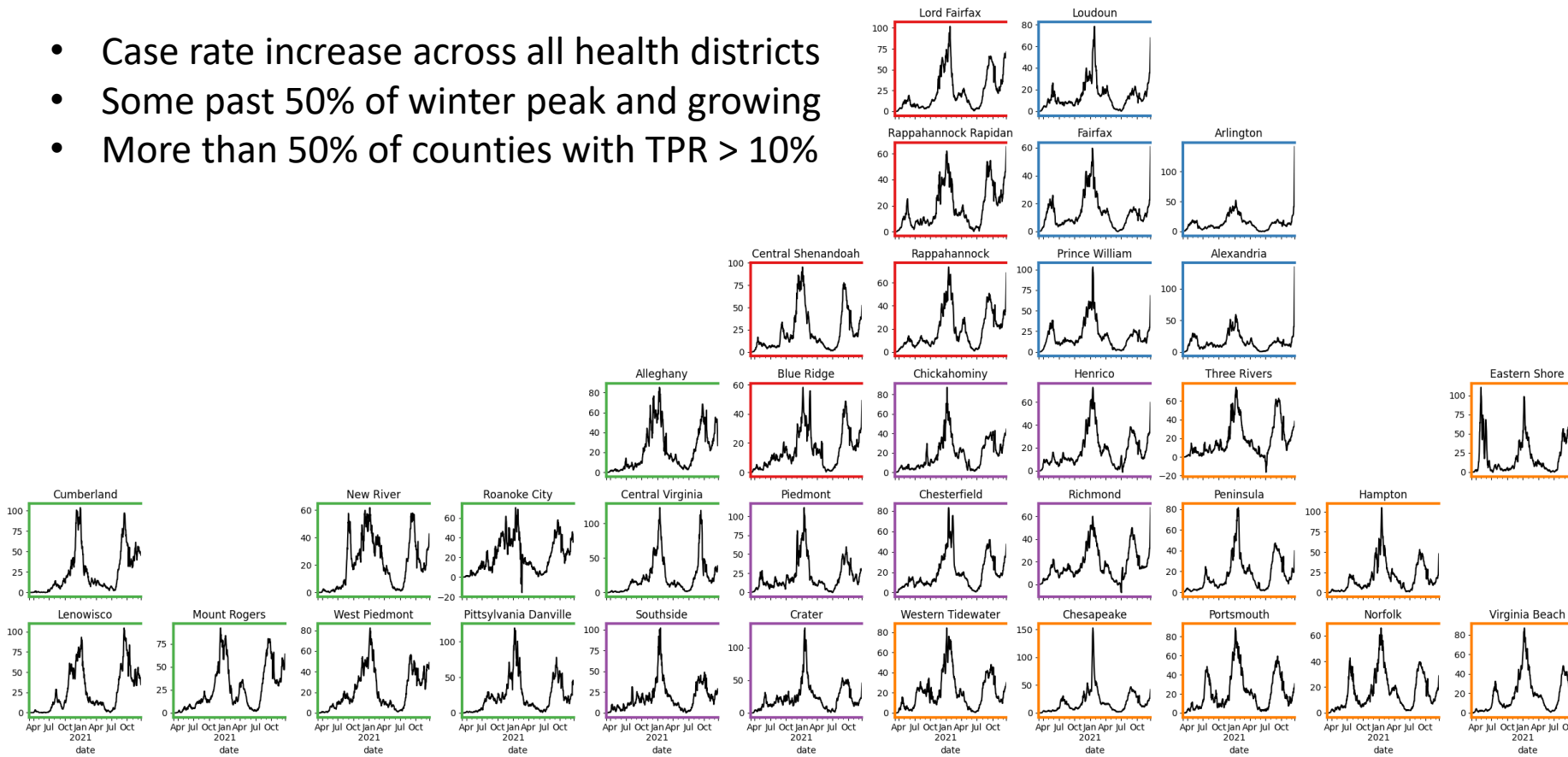


## County level RT-PCR test positivity

**Green:** <5.0% (or <20 tests in past 14 days)

**Yellow:** 5.0%-10.0% (or <500 tests and <2000 tests/100k and >10% positivity over 14 days)

**Red:** >10.0% (and not "Green" or "Yellow")



# District Trajectories

**Goal:** Define epochs of a Health District's COVID-19 incidence to characterize the current trajectory

**Method:** Find recent peak and use hockey stick fit to find inflection point afterwards, then use this period's slope to define the trajectory

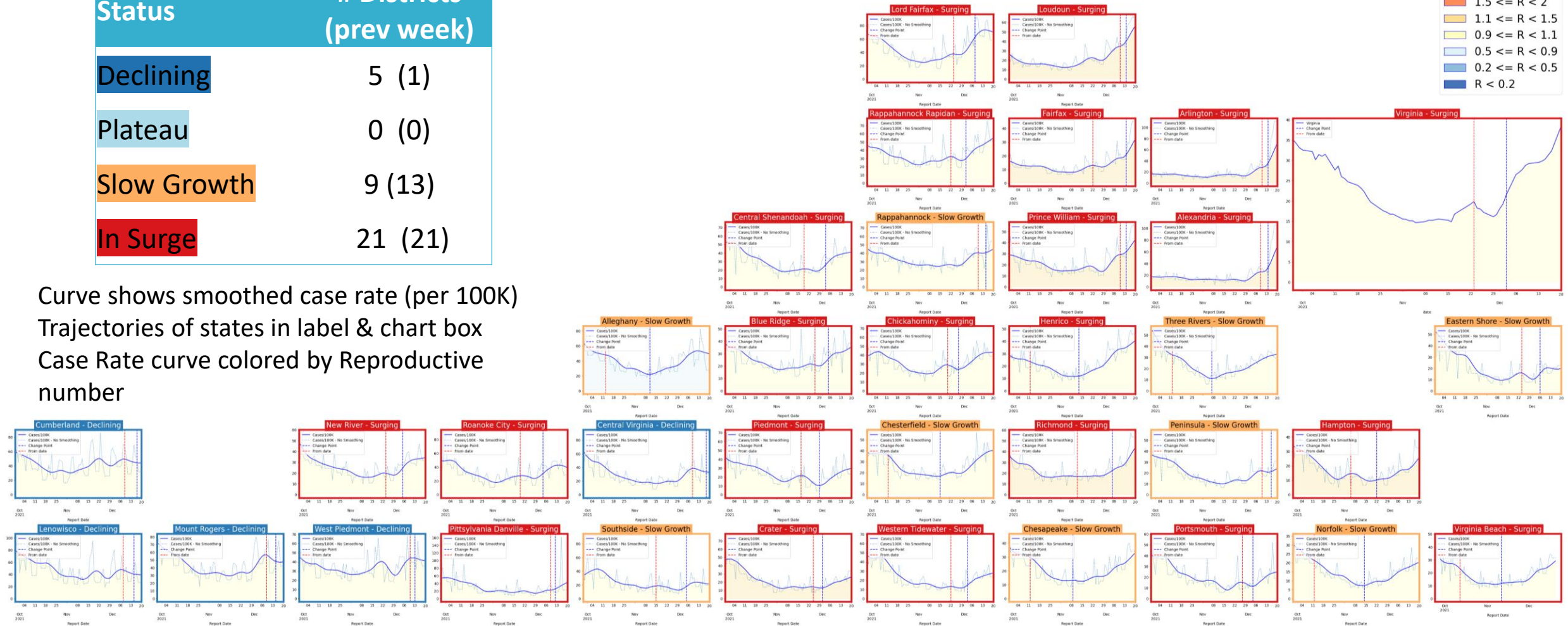
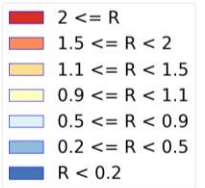


| Trajectory         | Description   | Weekly Case Rate (per 100K) bounds | # Districts (prev week) |
|--------------------|---|------------------------------------|-------------------------|
| <b>Declining</b>   | Sustained decreases following a recent peak                   | below -0.9                         | 5 (1)                   |
| <b>Plateau</b>     | Steady level with minimal trend up or down                    | above -0.9 and below 0.5           | 0 (0)                   |
| <b>Slow Growth</b> | Sustained growth not rapid enough to be considered a Surge    | above 0.5 and below 2.5            | 9 (13)                  |
| <b>In Surge</b>    | Currently experiencing sustained rapid and significant growth | 2.5 or greater                     | 21 (21)                 |

# District Trajectories – last 10 weeks

| Status      | # Districts<br>(prev week) |
|-------------|----------------------------|
| Declining   | 5 (1)                      |
| Plateau     | 0 (0)                      |
| Slow Growth | 9 (13)                     |
| In Surge    | 21 (21)                    |

Curve shows smoothed case rate (per 100K)  
Trajectories of states in label & chart box  
Case Rate curve colored by Reproductive number



# Estimating Daily Reproductive Number – Redistributed gap

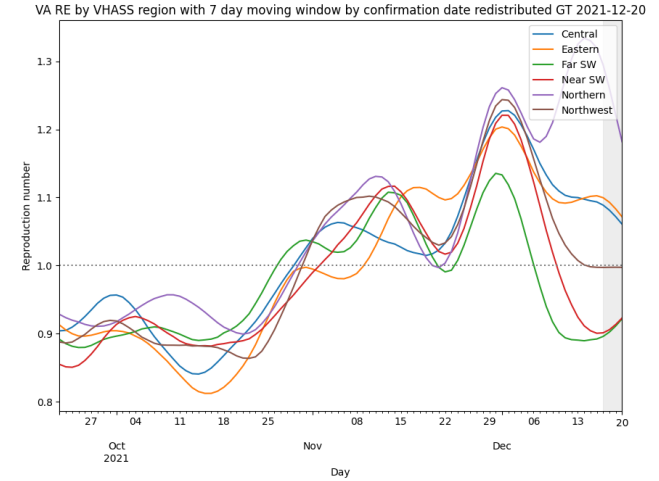
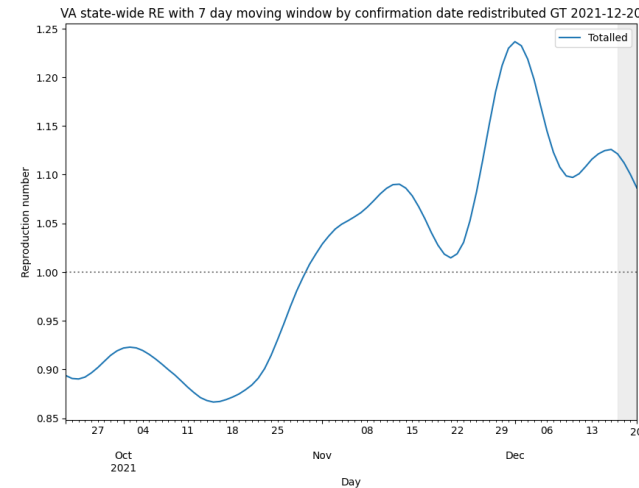
Dec 20<sup>th</sup> Estimates

| Region     | Date Confirmed<br>$R_e$ | Date Confirmed<br>Diff Last Week |
|------------|-------------------------|----------------------------------|
| State-wide | 1.116                   | -0.002                           |
| Central    | 1.059                   | 0.002                            |
| Eastern    | 1.072                   | 0.051                            |
| Far SW     | 0.919                   | 0.025                            |
| Near SW    | 0.923                   | -0.086                           |
| Northern   | 1.181                   | 0.149                            |
| Northwest  | 0.996                   | -0.056                           |

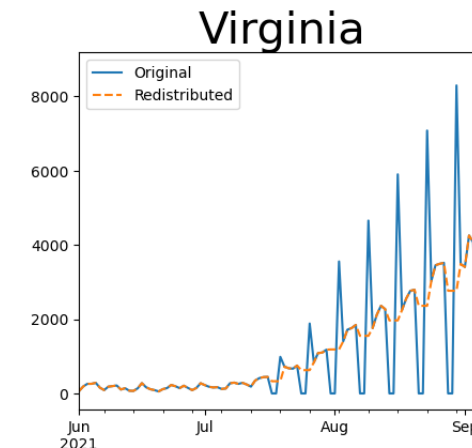
## Methodology

- Wallinga-Teunis method (EpiEstim<sup>1</sup>) for cases by confirmation date
- Serial interval: updated to discrete distribution from observations (mean=4.3, Flaxman et al, Nature 2020)
- Using Confirmation date since due to increasingly unstable estimates from onset date due to backfill

1. Anne Cori, Neil M. Ferguson, Christophe Fraser, Simon Cauchemez. A New Framework and Software to Estimate Time-Varying Reproduction Numbers During Epidemics. American Journal of Epidemiology, Volume 178, Issue 9, 1 November 2013, Pages 1505–1512, <https://doi.org/10.1093/aje/kwt133>



Skipping Weekend Reports & holidays biases estimates  
Redistributed “big” report day to fill in gaps, and then estimate R from “smoothed” time series

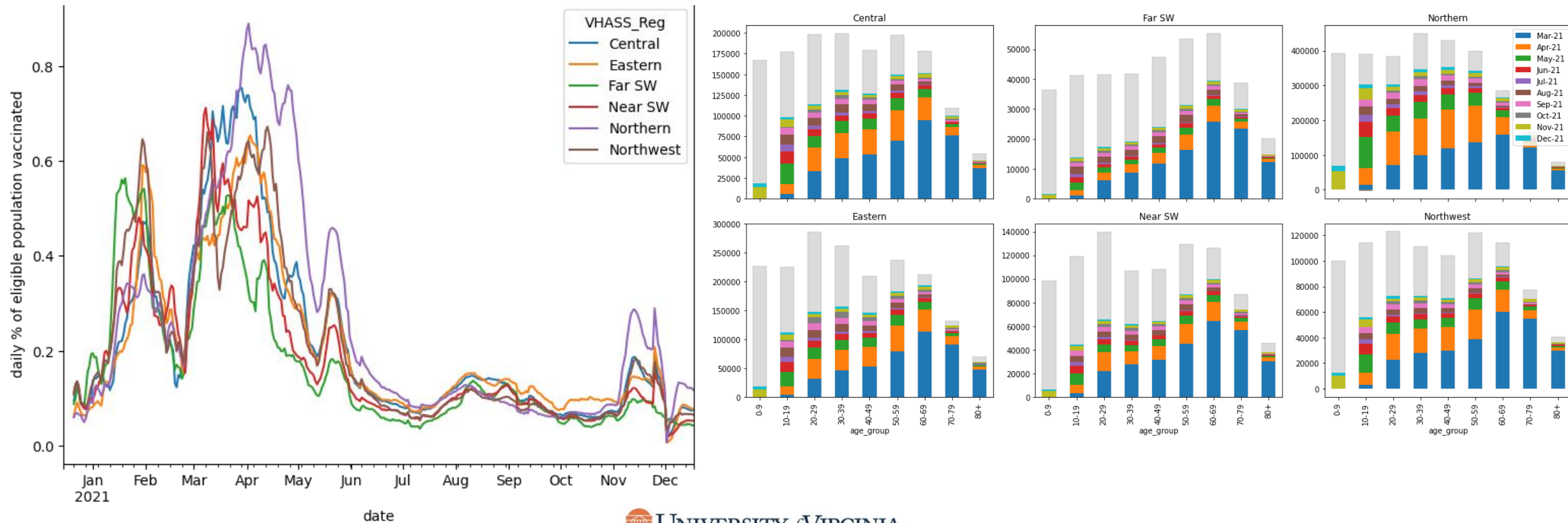




# Vaccination Administration Slow

## Regional Vaccine courses initiated per day (% eligible):

- Proportion eligible for first dose of vaccines across regions (in the ~0.1% or 100 per 100K a day)
- Age-specific proportions of population vaccinated show recent progress in younger ages

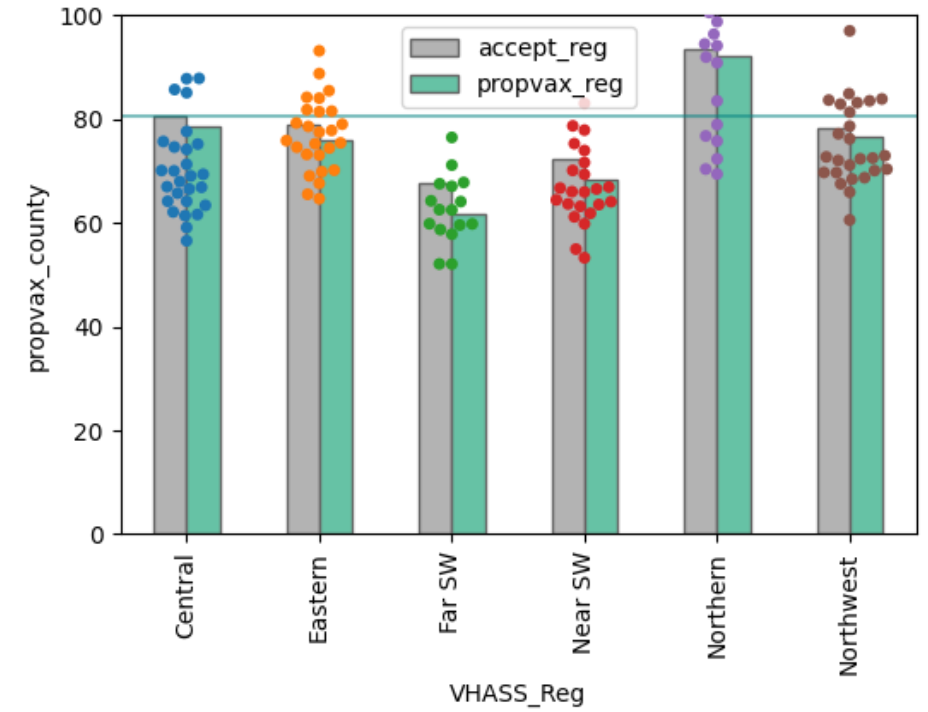


# Vaccination Acceptance by Region

## Corrections to surveys:

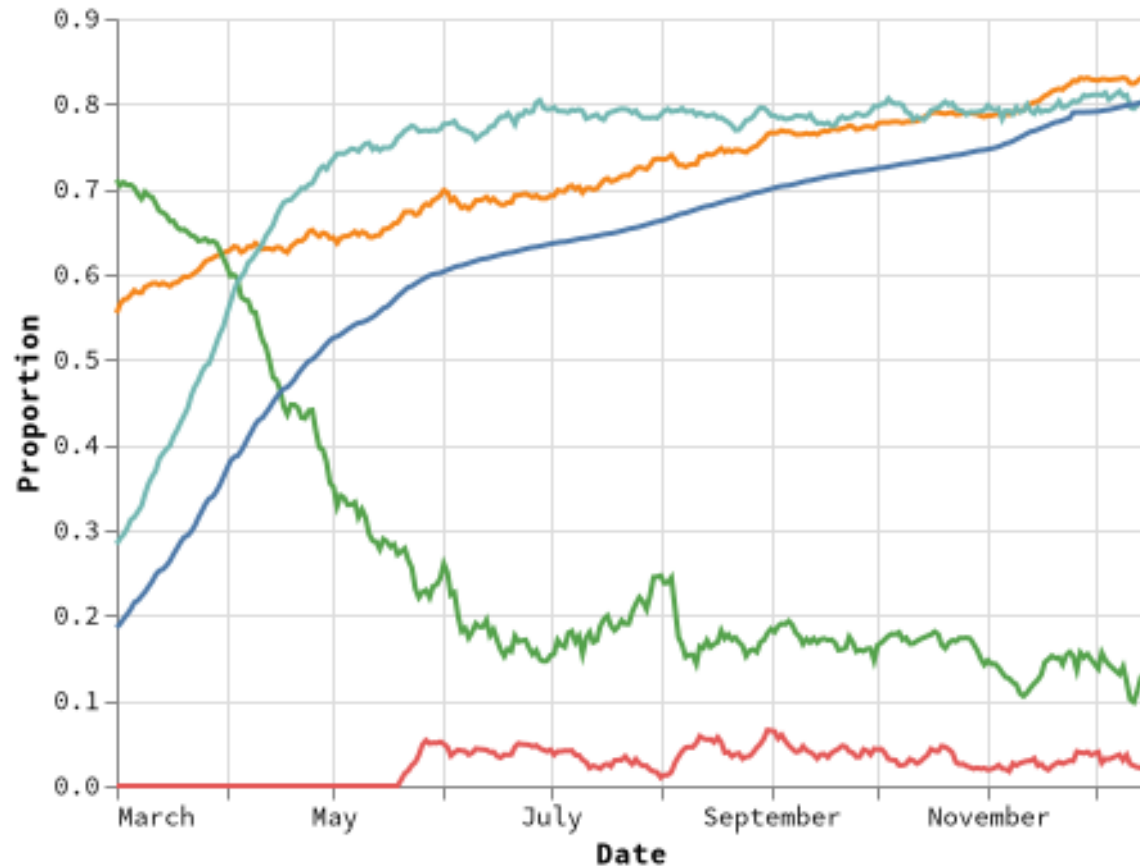
- Facebook administered survey is timely and broad, but biased by who accesses Facebook and answers the survey
- Correction approach:
  - Calculate an over-reporting fraction based on reported vaccinations compared to VDH administration data
  - Cross-validate coarse corrections against HPS survey at the state level and corrected in same manner

| Region          | COVIDcast accepting corrected | VDH proportion pop vaccinated |
|-----------------|-------------------------------|-------------------------------|
| Central         | 80%                           | 78%                           |
| Eastern         | 79%                           | 75%                           |
| Far SW          | 65%                           | 62%                           |
| Near SW         | 72%                           | 68%                           |
| Northern        | 94%                           | 92%                           |
| Northwest       | 78%                           | 76%                           |
| <b>Virginia</b> | <b>83%</b>                    | <b>80%</b>                    |



**Grey Bar:** Survey measured and corrected acceptance  
**Green Bar:** Proportion of eligible population administered a vaccine  
**Dots:** Proportion administered at least one dose for each county

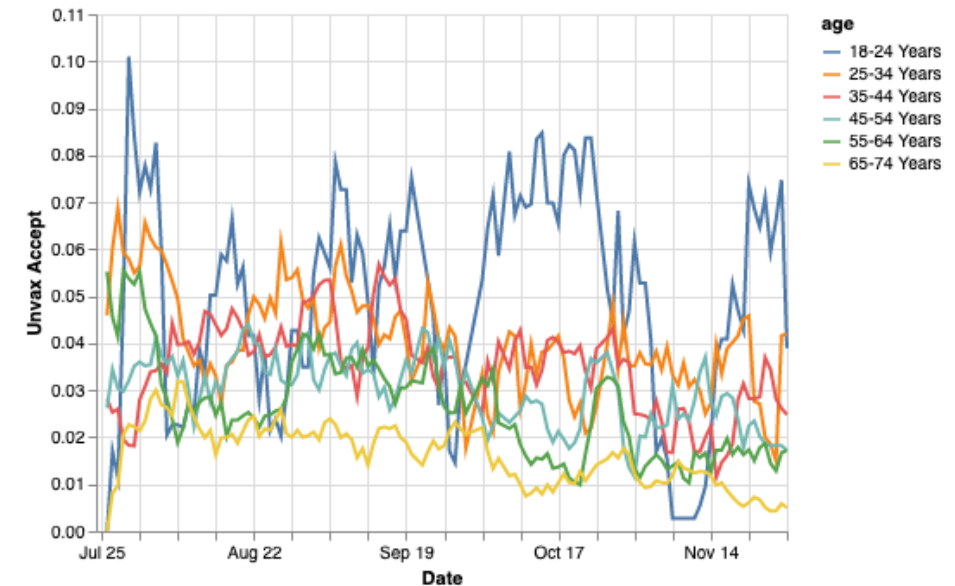
# Vaccine Acceptance Components over Time



## Vaccine Acceptance adjusted to include scheduled appointments

- Steady rise in acceptance over the past couple months
- Unvaccinated Acceptance shows ~20% of those who are unvaccinated are definitely or probably willing to be vaccinated
- Scheduled appointments for vaccination have increased through August but seem to be leveling off

## Acceptance Across Age groups among Unvaccinated



Data Source: <https://covidcast.cmu.edu>

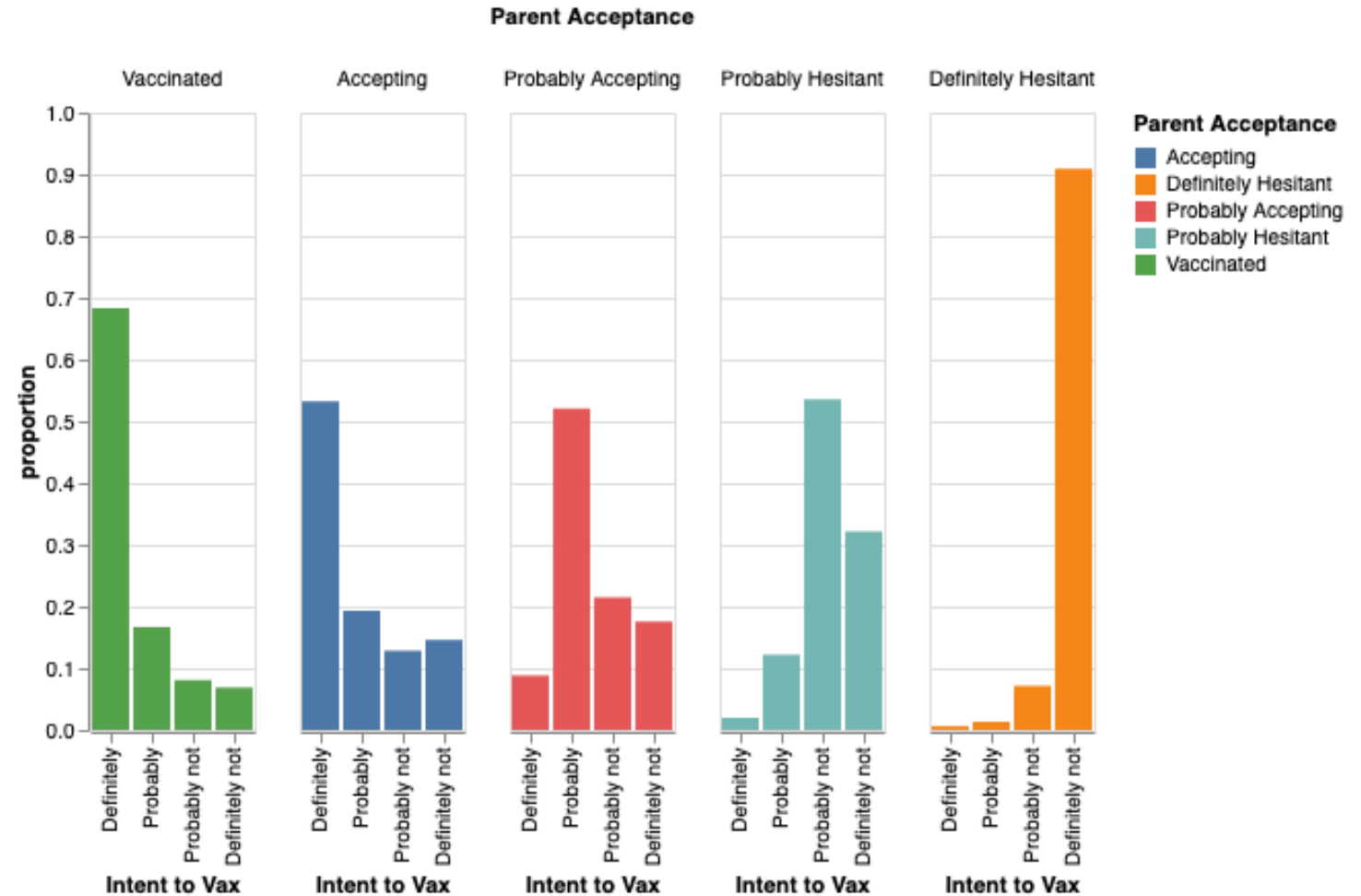
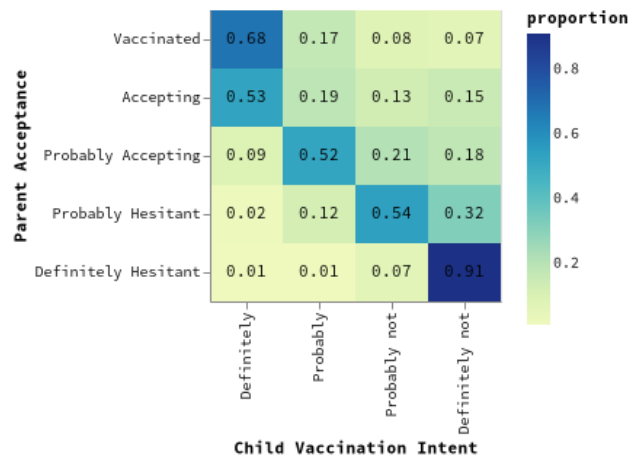
22-Dec-21



# Parental Intention to Vaccinate Children

## Parental Intention to Vaccinate Children lower than overall Acceptance

- Most willing (vaccinated) remain at ~70% definitely intending to vaccinate kids
- Intention strongly biased by the willingness of the parent, and skews towards unwillingness to vaccinate



Data Source: <https://covidcast.cmu.edu>

22-Dec-21

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# Mask Usage Stalls

**Self-reported mask usage has increased slightly to ~64% (mid 60s in previous months)**

- US and VA experienced similar small ticks up
- Mask wearing remains lower amongst unvaccinated especially among least willing to be vaccinated

*In the past 7 days, did you wear a mask **most or all of the time** in public?*

On Sun, Dec 19th 2021 the 7 day average of **People Wearing Masks (Last 7 Days)** was:

**64.3**

per 100 people

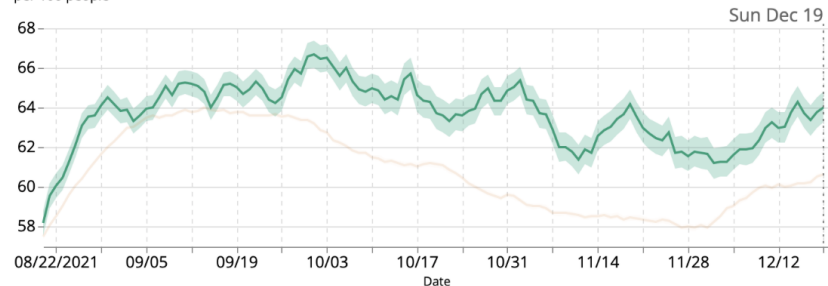
**+1.63%**

Relative change to 7 days ago



The indicator **People Wearing Masks (Last 7 Days)** was added in [Wave 8](#) of the Delphi survey published on Sun, Feb 7th 2021.

People Wearing Masks (Last 7 Days) in Virginia  
per 100 people



Delphi Group, [delphi.cmu.edu/covidcast](https://delphi.cmu.edu/covidcast)

☐ Include 0 in Y Axis ☐ Show All Dates

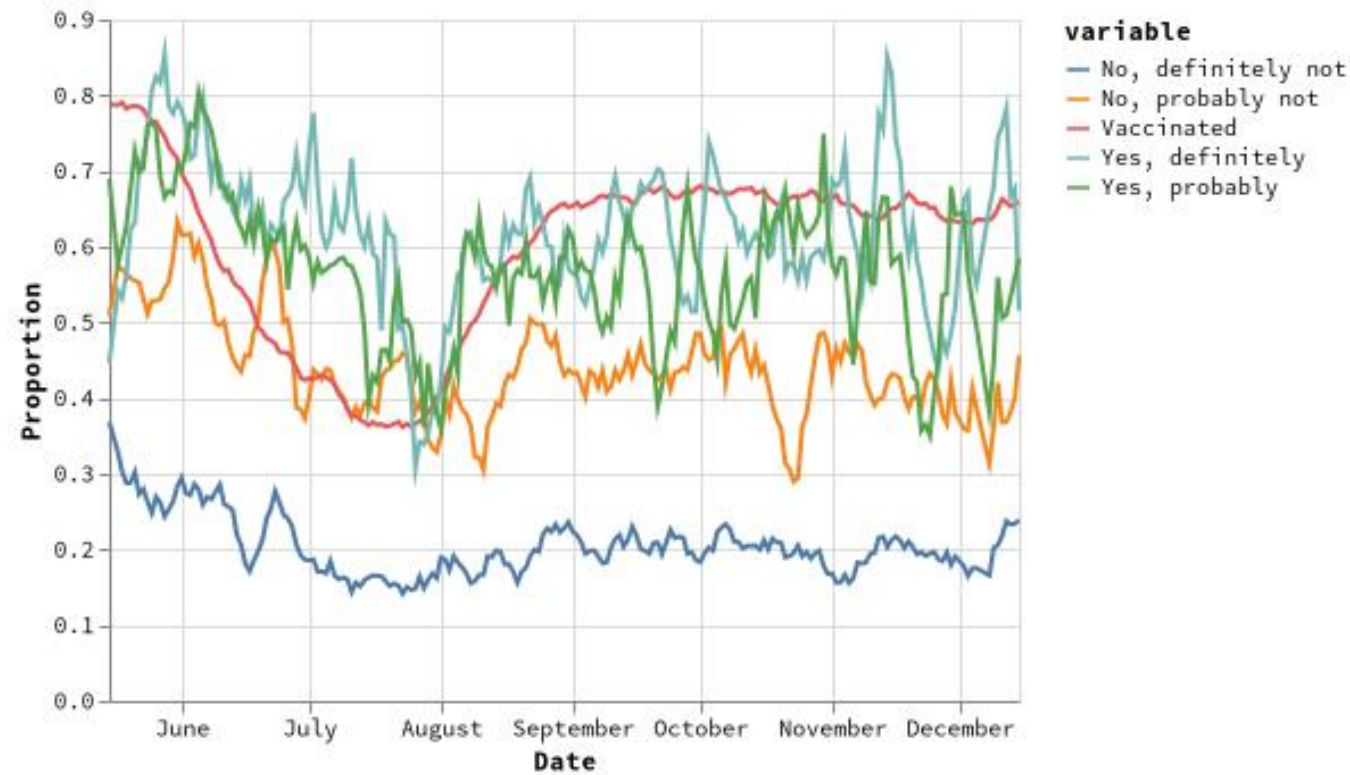
• Virginia

64.03 per 100

• United States

60.62 per 100

The 7-day average of "People Wearing Masks (Last 7 Days)" on 12/19/2021 is based on 4,003 samples. It was published on 12/21/2021.



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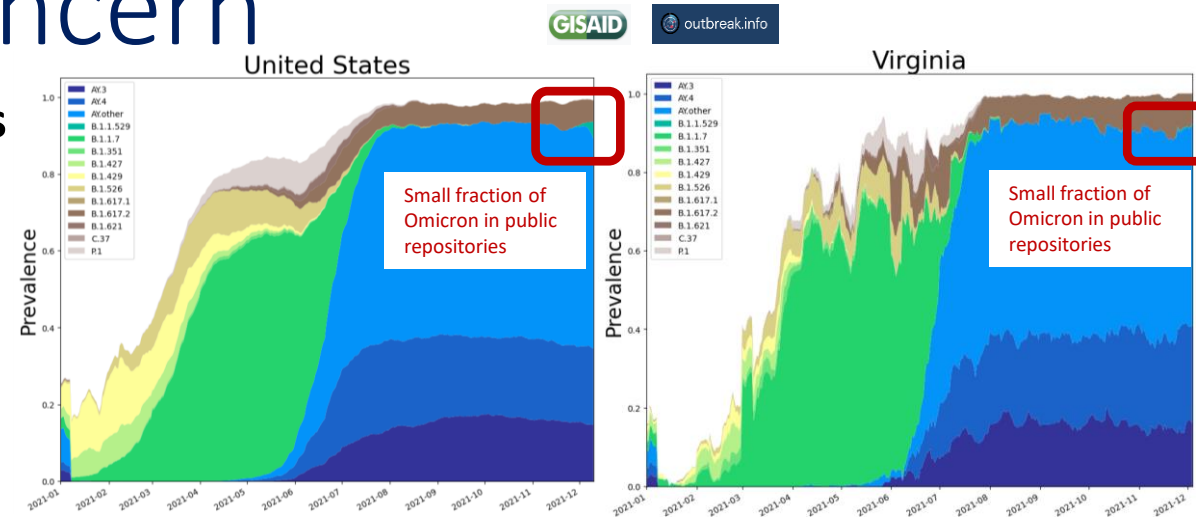
Data Source: <https://covidcast.cmu.edu>

# SARS-CoV2 Variants of Concern

Emerging new variants will alter the future trajectories of pandemic and have implications for future control

- Emerging variants can:
  - Increase transmissibility
  - Increase severity (more hospitalizations and/or deaths)
  - Limit immunity provided by prior infection and vaccinations
- Genomic surveillance remains very limited
  - Challenges ability to estimate impact in US to date and estimation of arrival and potential impact in future

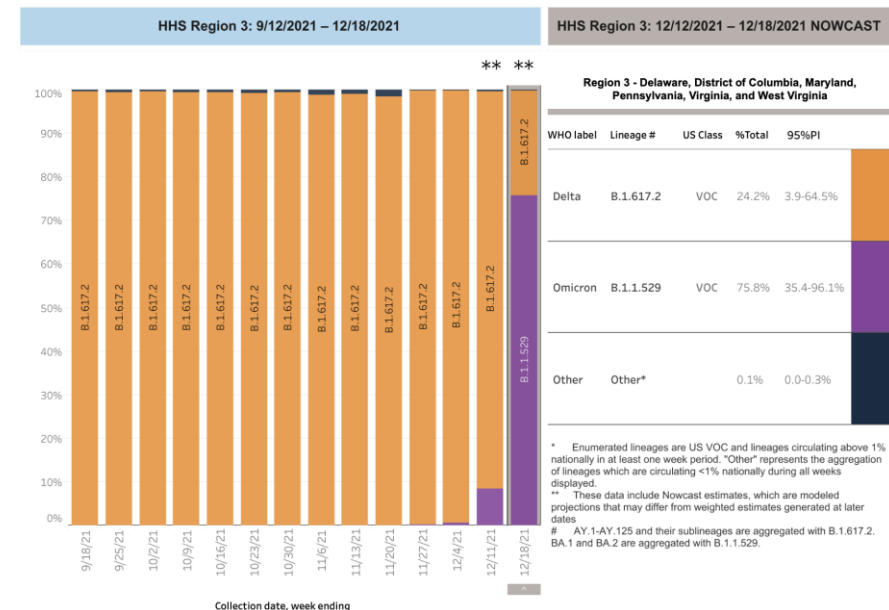
| WHO label | Pango lineage* | GISAID clade | Nextstrain clade | Additional amino acid changes monitored* | Earliest documented samples  | Date of designation                  |
|-----------|----------------|--------------|------------------|--|------------------------------|--------------------------------------|
| Alpha     | B.1.1.7        | GRY          | 20I (V1)         | +S:484K<br>+S:452R                       | United Kingdom, Sep-2020     | 18-Dec-2020                          |
| Beta      | B.1.351        | GH/501Y.V2   | 20H (V2)         | +S:L18F                                  | South Africa, May-2020       | 18-Dec-2020                          |
| Gamma     | P.1            | GR/501Y.V3   | 20J (V3)         | +S:681H                                  | Brazil, Nov-2020             | 11-Jan-2021                          |
| Delta     | B.1.617.2      | GI/478K.V1   | 21A, 21I, 21J    | +S:417N<br>+S:484K                       | India, Oct-2020              | VOI: 4-Apr-2021<br>VOC: 11-May-2021  |
| Omicron*  | B.1.1.529      | GRA          | 21K, 21L         | +R346K                                   | Multiple countries, Nov-2021 | VUM: 24-Nov-2021<br>VOC: 26-Nov-2021 |



**Omicron Prevalence**  
**SIGNIFICANTLY revised**  
**from last week**

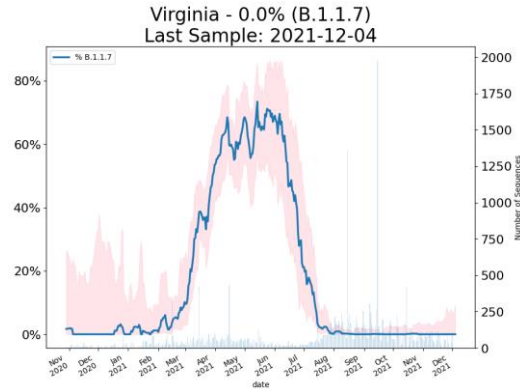
**Region 3:**  
8.4% to 76% in a week  
~3 doublings

**National:**  
12% to 73% in a week  
~ 3 doublings

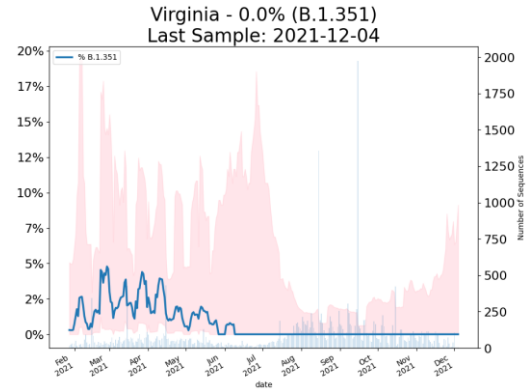


# SARS-CoV2 Variants of Concern

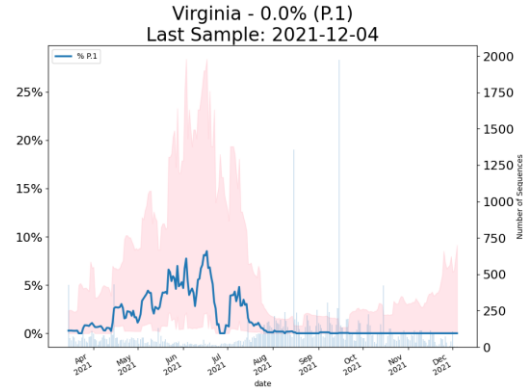
## Alpha $\alpha$ - Lineage B.1.1.7



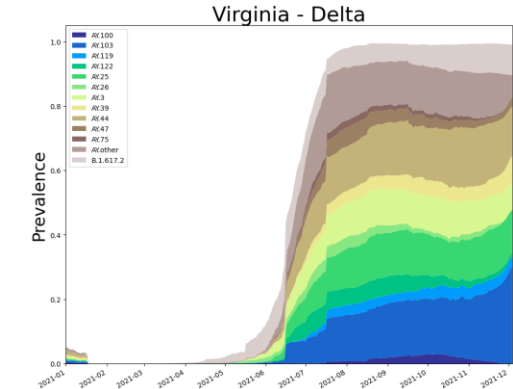
## Beta $\beta$ - Lineage B.1.351



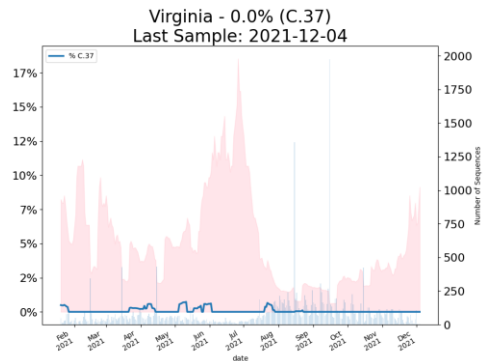
## Gamma $\gamma$ - Lineage P.1



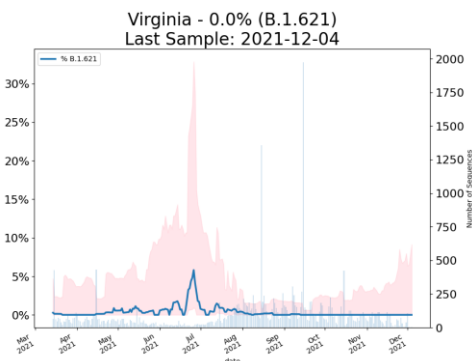
## Delta $\delta$ - Lineage B.1.617.2



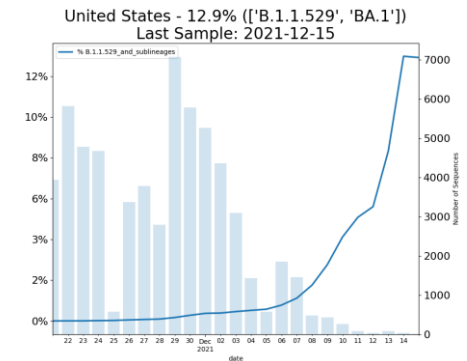
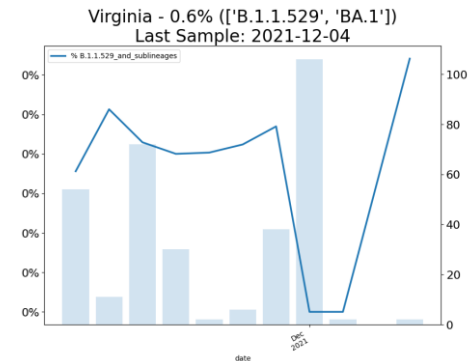
## Lambda $\lambda$ - Lineage C.37



## Mu $\mu$ - Lineage B.1.621



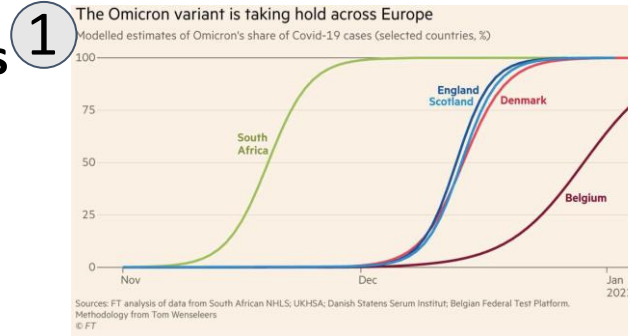
## Omicron $\omicron$ - Lineage B.1.1.529



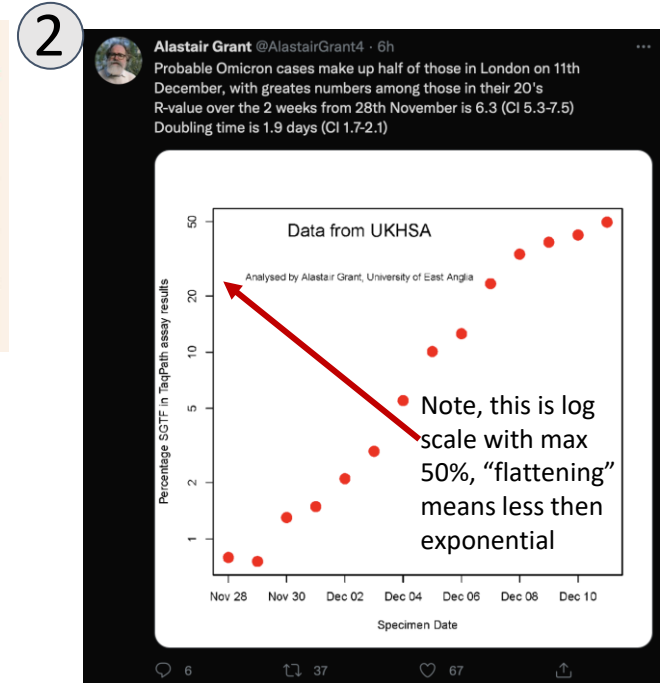
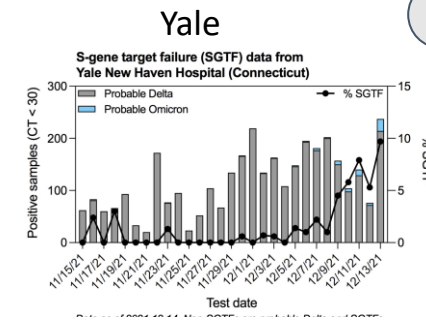
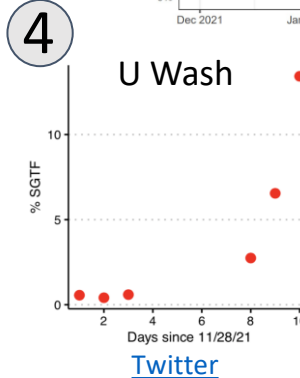
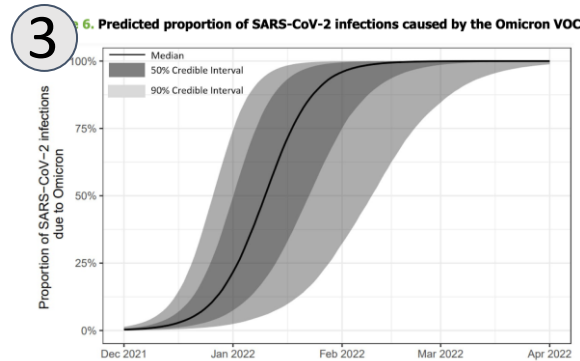
# Omicron – Prevalence and Growth

Omicron has become dominant fast and fuels case rate growth

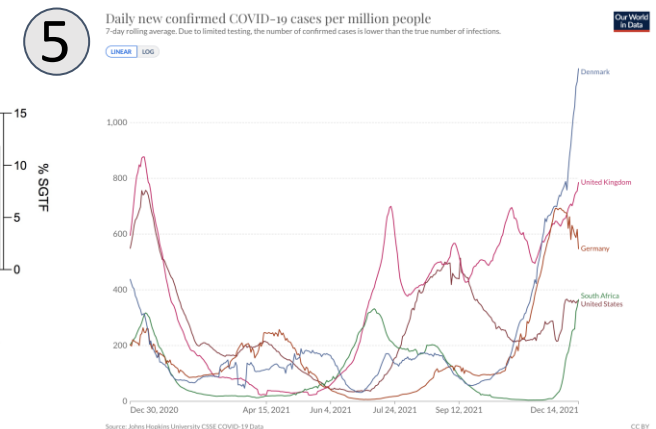
1. Experience of South Africa is being followed in several European nations ([Financial Times](#))
2. Omicron now dominant strain in UK (below 1% to 50% in 2 weeks), growth of SGTF% in UK remains exponential, with only slight slowing ([Twitter](#))
3. ECDC modeling estimates predominance in early 2022 for Europe ([ECDC](#) via [Twitter](#))
4. Univ Washington and Yale both report over 10% for recent SGTF%
5. Case Rate growth in countries with lots of Omicron has been more rapid than previous waves ([Our World in Data](#))



<https://www.ft.com/content/3c27c135-fdbc-4db7-8c7c-6e1f6c386235>



<https://twitter.com/AlastairGrant4/status/1471032314436235268>



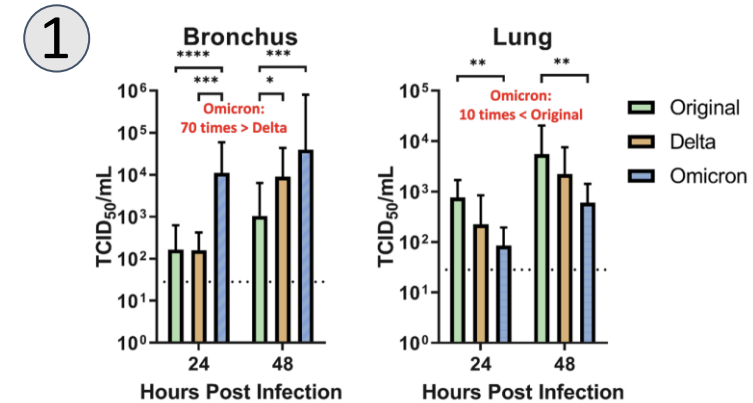
[Our World in Data](#)



# Omicron – Immune Evasion and Transmissibility

## Lab Studies find Omicron to be more transmissible and immune evading

1. Study from Hong Kong shows Omicron to be much more efficient at infecting Bronchus cells, though slightly less in Lung cells ([HKU](#))
2. In vitro study finds antibody neutralization of Omicron to be much stronger from serum from individuals with a 3<sup>rd</sup> dose of mRNA vaccine than just 2 doses ([non-peer reviewed preprint](#))
3. Initial analysis of Omicron mutations suggest there is minimal evasion against T-cell mediated immune response, thus T-cell mediated immunity should remain robust against Omicron, ie infection may occur but immune response will still occur ([BioArxiv](#))



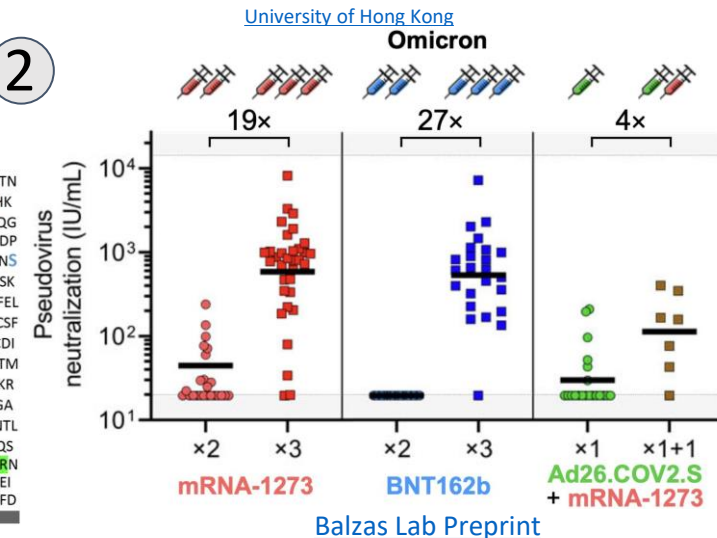
**3**

**Spike**

MFVFLVLLPVSSQCVNLTTRTQLPPAYTNSFTRGVVYPDKVFRSSVLSHTQDLFLPFFSNVTWFHAIHVSGTN  
 GTRFDNPVLPFNDGVYFAS<sup>TE</sup>SNIRGWIFGTTLDSTQSLVNNATNVVIVKCEFCNDPFLGVVYHK  
 NNKSWMESEFRVSSANNCTFEYVQQFLMDLEGKQGNFKNREFVKNIDGYFKIYKHTPI<sup>IN</sup>LVRLDLPQG  
 FSALEPLVDLPIGINITRFQTLALHRSYLTGPDSSSGWTAGAAAYVYG<sup>YLQPR</sup>TFLKYNENGTTITDAVDCALDP  
 LSETKCTLKSFVKEGIYQTSNFRVQPTESIVRFPNITNLCPE<sup>GE</sup>VFNATRFASVYAWNRKRISNCVADSVLYNS  
 ASFSTFKCYGVSPTKLNDLCTNVYADSFVIRGDEVRIAPGQGTG<sup>KI</sup>ADYNYKLPPDFTGCVIAWNSN<sup>NL</sup>DSK  
 VGG<sup>NY</sup>NLYRLFRKSNLK<sup>PF</sup>ERDISTEIQAG<sup>ST</sup>PCNGVEGFNCFYPLQSY<sup>GF</sup>QPTNGVGYPYRVVLSFEL  
 LHAPATVCGPKKSTNLVKNKCVNFNGLTGTGVLTESNKKFLPFQFGRDIAD<sup>TD</sup>AVRDPQTLEILDITPCSF  
 GGVSVITPGTNTSNQAVLYQ<sup>DN</sup>CTEVPVAIHADQLTPTWRVYSTGSNVFQTRAGCLIGAE<sup>HV</sup>NNSYECDI  
 PIGAGICASYQTQT<sup>NS</sup>PRRARSVASQSIAYTMSLGAENSVAYSNNIAIPTNFTISVTTEILPVSMTKTSDVCTM  
 YICGDSTECNLLQYGSFCTQL<sup>N</sup>RALTGIAVEQDKNTQEVFAQVKQIYKTPPIK<sup>DF</sup>GGFNFSQILPDPSKPSKR  
 SFIEDLLFNKVTLDAGFIKQYGDCLGDIARDLCAQKF<sup>NG</sup>LTVLPPL<sup>LD</sup>EMIAQYTSALLAGTITSGWTFGA  
 GAALQIPFAMQMAYRFNGIGVTQNVLYENQFNQFSAIGIKQDSLSSTASALGKLQDVVN<sup>Q</sup>NAQALNTL  
 VKQLSSN<sup>FG</sup>AISSVLNDIL<sup>SR</sup>LDKVEAEVQIDRLITGRLOSQTQYVYVQQLIRAAEIRASANLAATKMSCEVLGQS  
 KRVDFCGKGYHLMSPQS<sup>AP</sup>HGVVFLHVT<sup>TV</sup>PAQEKNTFTAPAICHGKAHPREGVFSVN<sup>GH</sup>WVFTQRN  
 FYEPQIITDNTFVSGNCDVIGVINYTVYDPLQPELDSFKEELDKYFKNHTSPDVLGDISGINASVNIQKEI  
 DRLNEVAKNLSLIDLQELGKYE<sup>QY</sup>KWPWY<sup>W</sup>LGFIAGLIAIVMTI<sup>ML</sup>CCMTSCCCLGKCCSCGSCCKFD

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[BioArxiv](#)



# Omicron – Severity

Assessing the severity of Omicron is challenging due to different risks by age, levels of immunity, types of vaccines used across the different populations affected by Omicron to date

1. [Study from Imperial College's MRC](#) found no evidence that Omicron cases were less likely to be admitted to hospital (and were much more likely to cause re-infection)
2. [Recent study from South Africa](#) found an adjusted OR of 0.3 for severe disease for Omicron compared to Delta, (severe disease defined as requiring ICU or similar advanced care)

S+ = Not Omicron  
S- = Proxy for Omicron

1

| Class              | Variable                 | S+     | S-    | log(OR)             | OR               | p-value |
|--------------------|--------------------------|--------|-------|---------------------|------------------|---------|
| Time               | Day                      | 208947 | 15087 | 0.43 (0.42-0.44)    | 1.54 (1.53-1.55) | <1e-6   |
|                    | London                   | 21585  | 5976  | 0                   | 1                | -       |
| NHS Region         | East of England          | 27986  | 2274  | -0.85 (-0.91--0.79) | 0.43 (0.4-0.46)  | <1e-6   |
|                    | Midlands                 | 41223  | 1645  | -1.53 (-1.59--1.46) | 0.22 (0.2-0.23)  | <1e-6   |
|                    | North East and Yorkshire | 45631  | 811   | -2.32 (-2.41--2.24) | 0.1 (0.09-0.11)  | <1e-6   |
|                    | North West               | 34726  | 1998  | -1.11 (-1.17--1.04) | 0.33 (0.31-0.35) | <1e-6   |
|                    | South East               | 27405  | 1985  | -0.85 (-0.92--0.79) | 0.43 (0.4-0.45)  | <1e-6   |
|                    | South West               | 10391  | 398   | -1.47 (-1.58--1.35) | 0.23 (0.21-0.26) | <1e-6   |
| Symptoms           | symptomatic              | 119284 | 8171  | 0                   | 1                | -       |
|                    | asymptomatic             | 89663  | 6916  | -0.02 (-0.06-0.02)  | 0.98 (0.95-1.02) | 0.4348  |
| Reinfection status | Not reinfection          | 206321 | 13586 | 0                   | 1                | -       |
|                    | Reinfection              | 2626   | 1501  | 1.88 (1.79-1.97)    | 6.55 (5.99-7.15) | <1e-6   |
| Hospital status    | No hospital attendance   | 207555 | 15063 | 0                   | 1                | -       |
|                    | Hospital attendance      | 1392   | 24    | -0.05 (-0.49-0.39)  | 0.95 (0.61-1.47) | 0.8275  |

[Imperial College Report](#)

## Early assessment of the clinical severity of the SARS-CoV-2 Omicron variant in South Africa

2

<sup>ID</sup> Nicole Wolter, Waasila Jassat, Sibongile Walaza, Richard Welch, Harry Moultrie, Michelle Groome, Daniel Gyamfi Amoako, Josie Everatt, Jinal N Bhiman, Cathrine Scheepers, Naume Tebeila, Nicola Chiwandire, Mignon du Plessis, Nevashan Govender, Arshad Ismail, Allison Glass, Koleka Mlisana, Wendy Stevens, Florette K Treurnicht, Zinhe Makatini, Nei-yuan Hsiao, Raveen Parboosing, Jeannette Wadula, Hannah Hussey, Mary-Ann Davies, Andrew Boule, Anne von Gottberg, <sup>ID</sup> Cheryl Cohen

doi: <https://doi.org/10.1101/2021.12.21.21268116>

**Table 3** Multivariable logistic regression analysis evaluating the association between S gene target failure (SGTF) infection during 1 October – 30 November 2021, compared to Delta variant infection during April – November 2021, and severe disease among hospitalized individuals with known outcome, South Africa<sup>a</sup> (N=1036)

|                    | Severe disease <sup>b</sup><br>n/N (%) | Adjusted odds ratio<br>(95% CI) | P-value |
|--------------------|--|---------------------------------|---------|
| SARS-CoV-2 variant | N=1037                                 |                                 |         |
| SGTF               | 57/244 (23)                            | 0.3 (0.2-0.5)                   | <0.001  |
| Delta              | 496/793 (63)                           | Ref                             | -       |

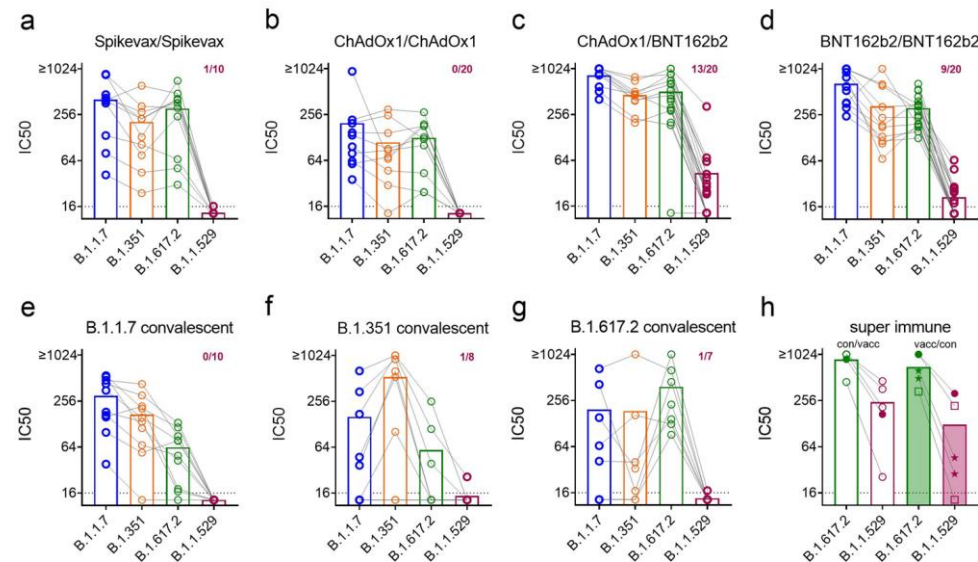
[MedArxiv](#)

| Indicator   | Risk assessment for SARS-CoV-2 variant: Omicron VOC-21NOV-01 (B.1.1.529) | Confidence level | Assessment and rationale   |
|---|--|------------------|--|
| Growth advantage  | Red  | High             | <b>Omicron is displaying a growth advantage over Delta</b><br>This assessment is based on analysis of UK data showing increased household transmission risk, increased secondary attack rates and substantially increased growth rates compared to Delta. Omicron continues to increase as a proportion of UK cases and is now predominant in some regions of England. This growth advantage is also apparent in other countries with equivalent surveillance. The observed growth advantage may be due to immune evasion or transmissibility. Although we now have high confidence in a component of immune evasion, the very high growth rate and laboratory findings suggest that an increase in transmissibility may also be contributing. |
| Transmissibility  | Amber  | Low              | <b>Omicron is at least as transmissible as Delta</b><br>Increased transmissibility compared to Delta is biologically plausible with the presence of furin cleavage site and nucleocapsid changes associated in vitro with advantages for replication, as well as extensive changes to the RBD. Structural modelling suggests that the mutations present may increase human ACE2 binding affinity to a much greater extent than that seen for any other variant. Early laboratory data suggest more efficient cell entry and replication in bronchial cells in vitro. However, there is no clear epidemiological demonstration of transmissibility as distinct from other contributors to growth advantage.                                     |
| Immune evasion (including natural and vaccine derived immunity) | Red  | High             | <b>Omicron displays a reduction in immune protection against infection (NO data regarding severe disease)</b><br>This assessment is now based on neutralisation data from multiple laboratories, assessment of real world vaccine effectiveness in the UK and an observed increase in the risk of reinfection with Omicron. There are insufficient data to make any assessment of protection against severe disease.   |
| Infection severity  |  |                  | <b>Insufficient data</b><br>There are insufficient data to fully assess severity, which is expected in the early period of emergence of a new variant. However, on the data available in the UK, there is no signal that supports a difference in the intrinsic virulence of the Omicron virus compared to Delta.  |

A separate report will be published concerning impact on therapeutics in due course.

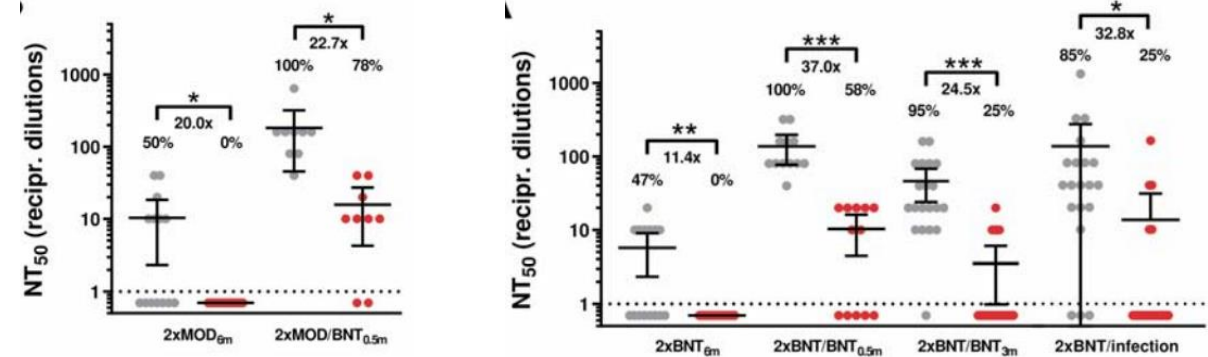
\* Refer to scale and confidence gradient slide

[https://twitter.com/dr\\_d\\_robertson/status/1468657221085147137](https://twitter.com/dr_d_robertson/status/1468657221085147137)



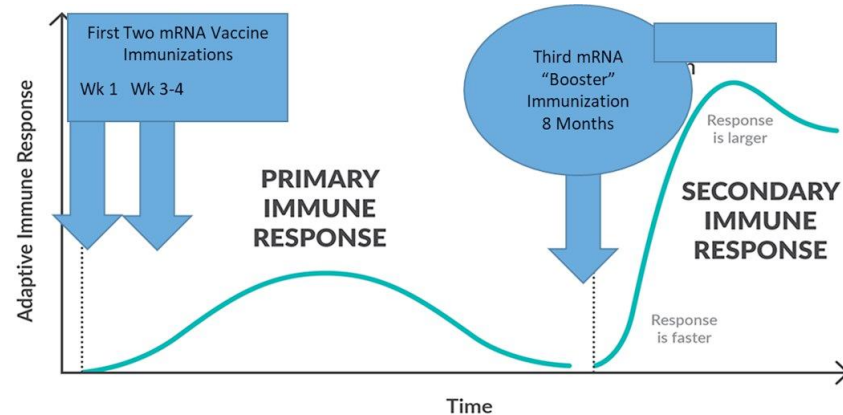
Kimpel, Medical University of Innsbruck  
<https://twitter.com/janinekimpe/status/1468700628922904591>

Omicron Red  
Delta grey



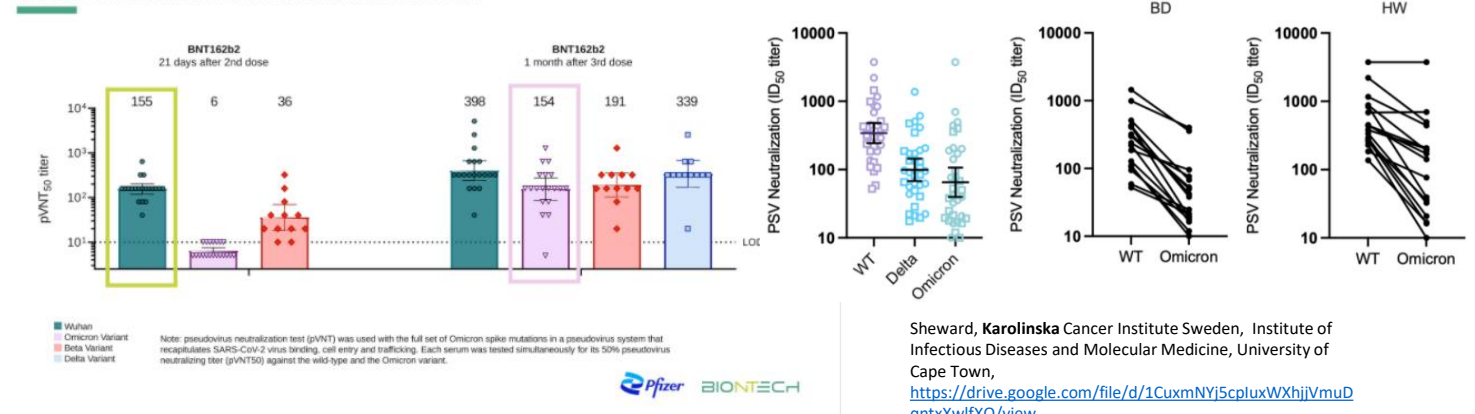
Ciesek, Institute for Medical Virology, University Hospital Frankfurt

<https://www.medrxiv.org/content/10.1101/2021.12.07.21267432v1.full.pdf>

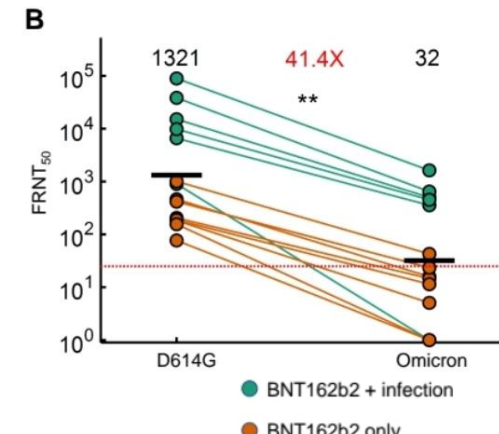


<https://twitter.com/peterhotez/status/1468712583146487812>

### Three doses of BNT162b2 neutralize Omicron



Pfizer study  
<https://twitter.com/EricTopol/status/1468593675747794946>



Signal, Max Planck Institute for Infection Biology (lab in Durban) (SA)  
<https://twitter.com/sigallab/status/1468325159501287434>

Sheward, Karolinska Cancer Institute Sweden, Institute of Infectious Diseases and Molecular Medicine, University of Cape Town,  
<https://drive.google.com/file/d/1CuxmNYj5cpluxWXhijVmuDqntxXwifXQ/view>

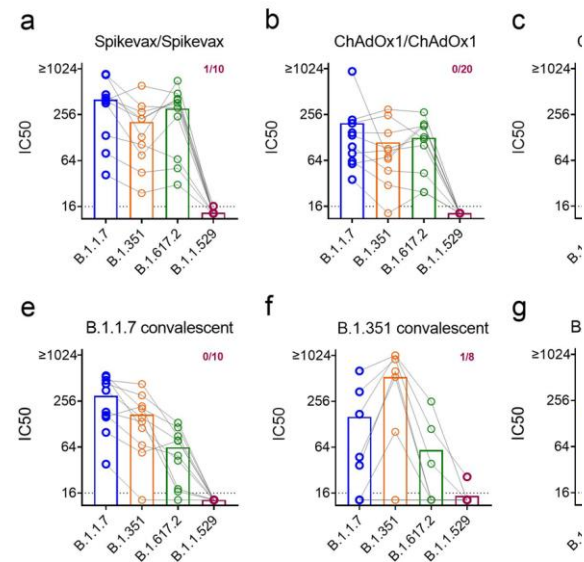


| Indicator   | Risk assessment for SARS-CoV-2 variant: Omicron VOC-2 | Confidence level | Assessment and rationale   |
|---|---|------------------|--|
| Growth advantage  | Red   | High             | Omicron is displaying a growth advantage. This assessment is based on transmission risk, increased spread compared to Delta. Omicron is predominant in some regions countries with equivalent surveillance immune evasion or transmissibility of immune evasion, the very high increase in transmissibility may increase the risk of reinfection with Omicron. |
| Transmissibility  | Amber   | Low              | Omicron is at least as transmissible as Delta. Increased transmissibility conferred by furin cleavage site and nucleocapsid replication, as well as extensive mutations present may increase the risk of reinfection with Omicron in bronchial cells in demonstration of transmissibility.   |
| Immune evasion (including natural and vaccine derived immunity) | Red   | High             | Omicron displays a reduction in neutralization regarding severe disease. This assessment is now based on assessment of real world vaccine risk of reinfection with Omicron protection against severe disease.  |
| Infection severity  |   |                  | Insufficient data. There are insufficient data to assess the emergence of a new variant that supports a difference in the risk of severe disease.  |

A separate report will be published concerning impact on therapeutics in due course.

\* Refer to scale and confidence gradient slide.

[https://twitter.com/dr\\_d\\_robertson/status/1468657221085147137](https://twitter.com/dr_d_robertson/status/1468657221085147137)

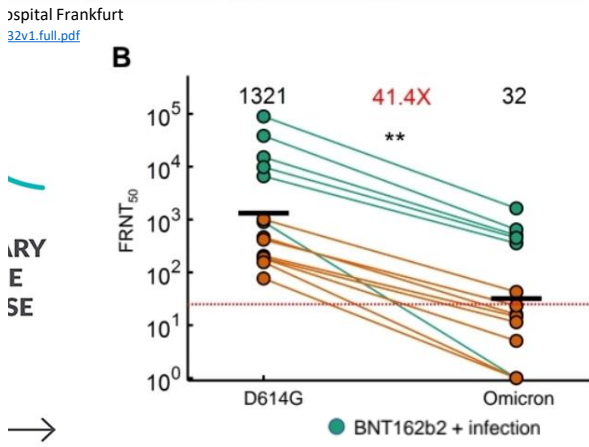
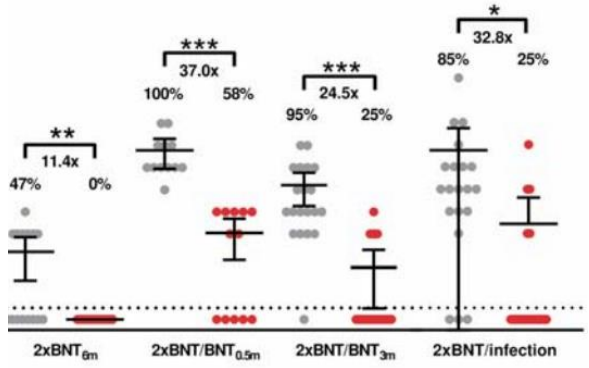


Kimpel, Medical University of Innsbruck  
<https://twitter.com/janinekimpel/status/1468700628922904591>

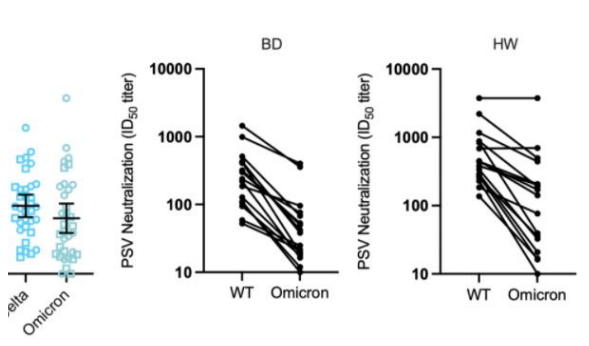
| Study       | Virus        | Sera   | N           | Fold reduction                      | Against ... | Remarks  | Link  |
|-------------|--------------|--|-------------|-------------------------------------|-------------|--|---|
| Sigal       | Live         | Pfizer (2 <sup>nd</sup> shot)  | 6           | 40x                                 | W           | Full manuscript available. Note that Omicron here had R346K                                    | <a href="https://bit.ly/omicron2">https://bit.ly/omicron2</a> |
|             |              | Recovered + Pfizer (2 <sup>nd</sup> shot)                                | 6           |                                     |             |  |   |
| Karolinska  | Pseudo lenti | Random Donors  | 17          | 7x                                  | W           | Full manuscript available. The details about the vaccination type of the donors are incomplete | <a href="https://bit.ly/omicron1">https://bit.ly/omicron1</a> |
|             |              | Infected   | 17          | 5x                                  |             |  |   |
| Ciesek      | Live         | Pfizer (2 <sup>nd</sup> shot)  | >5          | >10x (wipes out everyone below LoD) | D           | The number of people in each condition is unknown. The study describes more conditions         | <a href="https://bit.ly/omicron3">https://bit.ly/omicron3</a> |
|             |              | Pfizer (3 <sup>rd</sup> shot)  | >5          | ~30x                                |             |  |   |
|             |              | Pfizer (2 <sup>nd</sup> shot+ infected)                                  | >5          | 30x                                 |             |  |   |
| Pfizer      | Pseudo       | Pfizer (3 <sup>rd</sup> shot)  | 18(o) 12(Δ) | 2.5x                                | D           | A press release.   | <a href="https://bit.ly/omicron4">https://bit.ly/omicron4</a> |
| Kimpel      | Live         | Pfizer (2 <sup>nd</sup> shot)  | >10         | ~16x                                | D           | Full manuscript is now available   | <a href="https://bit.ly/omicron5">https://bit.ly/omicron5</a> |
| Zhang       | Pseudo       | Convalescent   | 28          | 8x                                  | W           | A full manuscript  | <a href="https://bit.ly/omicron6">https://bit.ly/omicron6</a> |
| Schmidt     | Pseudo       | Pfizer (2 <sup>nd</sup> shot) and Pfizer (3 <sup>rd</sup> shot after 1m) | 54 (Pfizer) | 16x                                 | W           | A full manuscript. Spike includes R683G. The study describes more conditions                   | <a href="https://bit.ly/omicron7">https://bit.ly/omicron7</a> |
| Israeli MoH | Live         | Pfizer (2 <sup>nd</sup> shot)  | ?           | 20x                                 | W           | Just a tweet from TV for now   | <a href="https://bit.ly/omicron8">https://bit.ly/omicron8</a> |
|             |              | Pfizer (3 <sup>rd</sup> shot 1M)   | ?           | 9x                                  | W           |  |   |

W: Ancestral; D: Delta

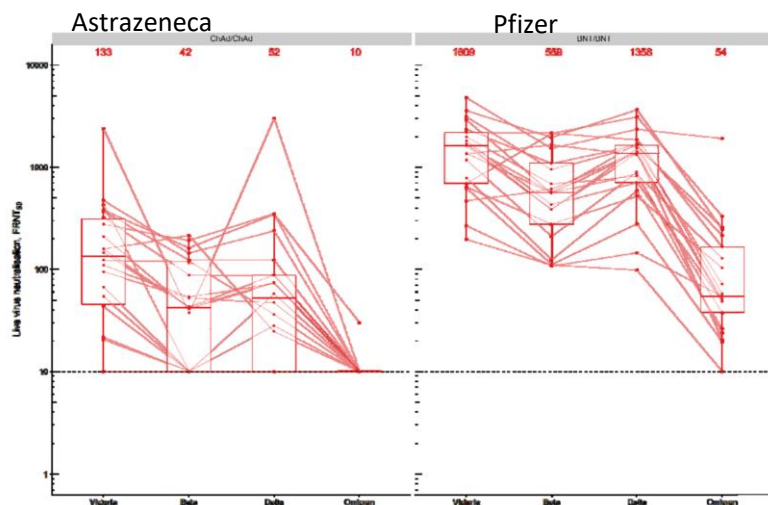
By @erlichya



Sigal, Max Planck Institute for Infection Biology (lab in Durban, SA)  
<https://twitter.com/sigallab/status/1468325159501287434>



Sheward, Karolinska Cancer Institute Sweden, Institute of Infectious Diseases and Molecular Medicine, University of Cape Town,  
<https://drive.google.com/file/d/1CuxmNYj5cpluxWXhijVmuDqntxXwifXQ/view>



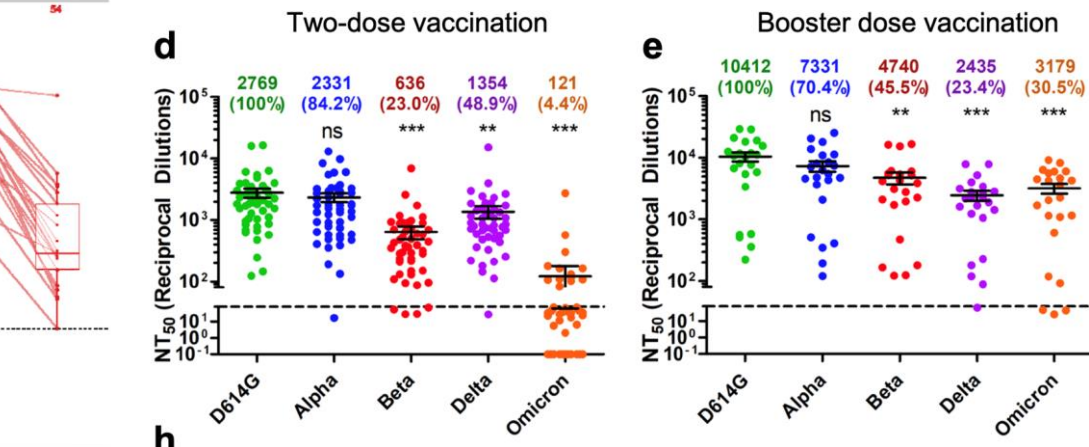
Oxford: Shows reduced live virus neutralisation of SARS-CoV-2 Omicron-B.1.1.529 variant by post immunisation serum  
<https://www.medrxiv.org/content/10.1101/2021.12.10.21267534v1.full.pdf>

**b**

|           | D614G | Alpha | Gamma | Beta  | Delta | Omicron |
|-----------|-------|-------|-------|-------|-------|---------|
| LY-CoV555 | 0.013 | 0.008 | >10   | >10   | >10   | >10     |
| LY-CoV016 | 0.032 | 1.707 | >10   | >10   | 0.024 | >10     |
| REGN10933 | 0.005 | 0.007 | 0.055 | 0.098 | 0.003 | >10     |
| REGN10987 | 0.005 | 0.003 | 0.003 | 0.002 | 0.005 | >10     |
| AZD8895   | 0.001 | 0.002 | 0.012 | 0.014 | 0.002 | 6.860   |
| AZD1061   | 0.001 | 0.001 | 0.002 | 0.003 | 0.004 | >10     |
| VIR-7831  | 0.058 | 0.080 | 0.066 | 0.050 | 0.073 | 0.181   |
| BR11-196  | 0.053 | 0.031 | 0.041 | 0.030 | 0.042 | 7.258   |
| DXP-604   | 0.010 | 0.007 | 0.005 | 0.065 | 0.016 | 0.287   |

IC50 (ng/μL)  
 0 5 10

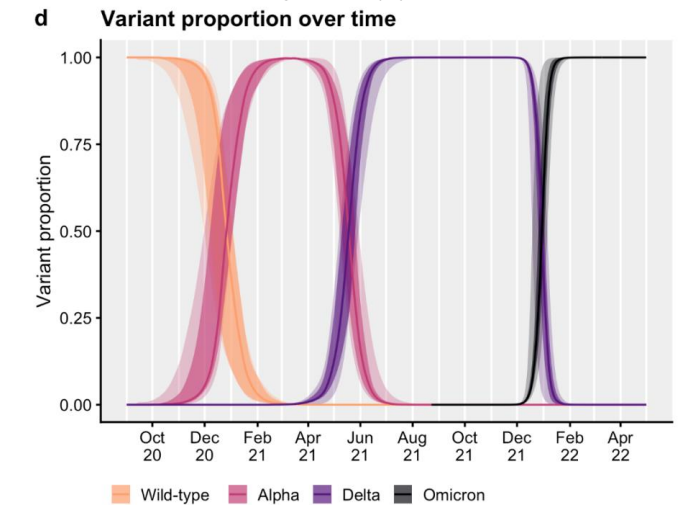
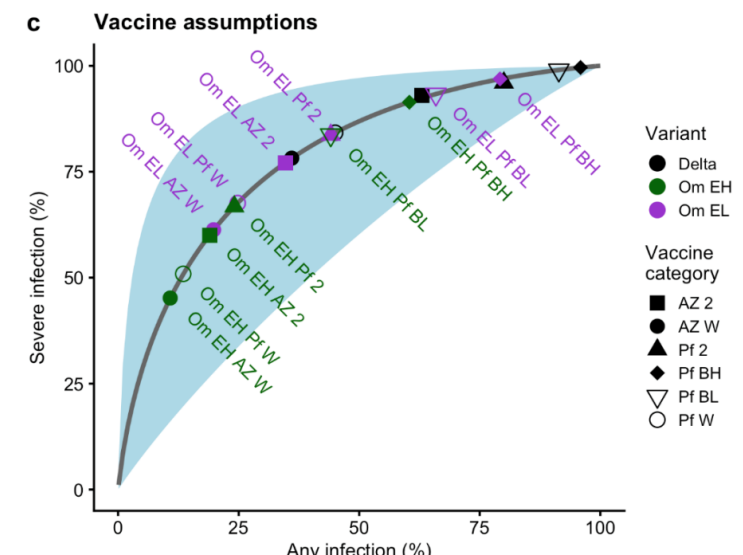
Beijing: B.1.1.529 escapes the majority of SARS-CoV-2 neutralizing antibodies of 2 diverse epitopes. 247 human anti-RBD NAb identified from SARS41 CoV/SARS-CoV-2 convalescents and vaccinees clustered into six epitope groups (A-F). Panel b shows antibody therapeutic neutralization performance. Panel C shows Nabs Groups E & F are the most resilient wrt Omicron neutralization but they are also the most rare.  
<https://www.medrxiv.org/content/10.1101/2021.12.10.21267534v1.full.pdf>



Ohio State: Findings suggest dual immune evasion strategies for Omicron, due to altered epitopes and reduced exposure of the S receptor binding domain, coupled with enhanced transmissibility due to enhanced S protein stability.  
<https://www.biorxiv.org/content/10.1101/2021.12.16.472934v1.article-info>

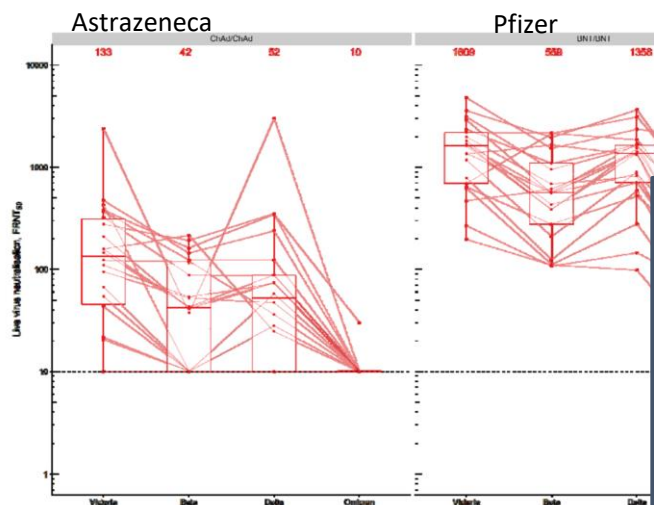
**c**

| Antibody Epitope Group | BD55-3152 | BD55-5319 | BD55-5386 | BD55-5300 | BD55-3372 | BD55-3500 |
|------------------------|-----------|-----------|-----------|-----------|-----------|-----------|
| Clade 1b               |           |           |           |           |           |           |
| SARS-CoV-1             | 3.462     | 3.447     | 3.417     | 3.405     | 3.467     | 3.476     |
| RaTG13                 | 0.353     | 0.135     | 0.086     | 0.193     | 0.061     | 3.258     |
| Pangolin-GD            | 1.132     | 2.657     | 0.421     | 3.526     | 3.493     | 3.57      |
| Pangolin-GX            | 0.996     | 0.812     | 1.95      | 0.709     | 2.889     | 3.403     |
| Clade 1a               |           |           |           |           |           |           |
| SARS-CoV-1             | 3.598     | 3.523     | 3.473     | 3.531     | 3.557     | 3.457     |
| PC4-127                | 3.458     | 3.495     | 3.468     | 3.464     | 3.413     | 2.984     |
| SARS-CoV-1             | 2.409     | 1.976     | 3.336     | 3.588     | 3.515     | 3.603     |
| Sin852                 |           |           |           |           |           |           |
| WIV1                   |           |           |           |           |           |           |
| Clade 3                |           |           |           |           |           |           |
| BM48-31                | 0.133     | 3.424     | 0.115     | 3.483     | 3.515     | 3.476     |
| Clade 2                |           |           |           |           |           |           |
| YN2013                 | 0.168     | 0.244     | 0.074     | 0.601     | 0.072     | 2.683     |



Edmunds, Davies et. al: We use the relationship between mean neutralisation titre and protective efficacy from Khoury et al. (7) to arrive at assumptions for vaccine efficacy against infection with Omicron, given each drop in neutralisation. We then use Khoury et al.'s modelled relationship between efficacy against any infection and efficacy against severe infection to generate vaccine effectiveness estimates against severe outcomes  
[https://cmmid.github.io/topics/covid19/reports/omicron\\_england/report\\_11\\_dec\\_2021.pdf](https://cmmid.github.io/topics/covid19/reports/omicron_england/report_11_dec_2021.pdf)





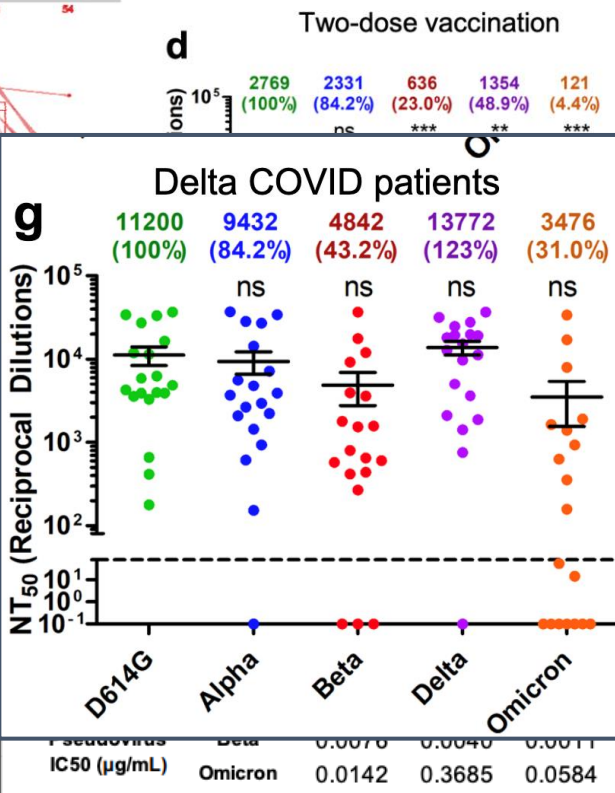
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**b**

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| REGN10987 | 0.005 | 0.003 | 0.003 | 0.002 | 0.005 | >10     |
| AZD8895   | 0.001 | 0.002 | 0.012 | 0.014 | 0.002 | 6.860   |
| AZD1061   | 0.001 | 0.001 | 0.002 | 0.003 | 0.004 | >10     |
| VIR-7831  | 0.058 | 0.080 | 0.066 | 0.050 | 0.073 | 0.181   |
| BR11-196  | 0.053 | 0.031 | 0.041 | 0.030 | 0.042 | 7.258   |
| DXP-604   | 0.010 | 0.007 | 0.005 | 0.065 | 0.016 | 0.287   |

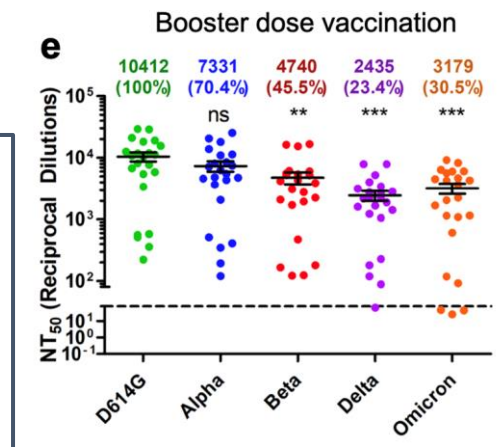
IC50 (ng/μL)  
 0 5 10

Beijing: B.1.1.529 escapes the majority of SARS-CoV-2 neutralizing antibodies of 2 diverse epitopes. 247 human anti-RBD NAb identified from SARS41 CoV/SARS-CoV-2 convalescents and vaccinees clustered into six epitope groups (A-F). Panel b shows antibody therapeutic neutralization performance. Panel c shows Nabs Groups E & F are the most resilient wrt Omicron neutralization but they are also the most rare.  
<https://www.medrxiv.org/content/10.1101/2021.12.10.21267534v1.full.pdf>

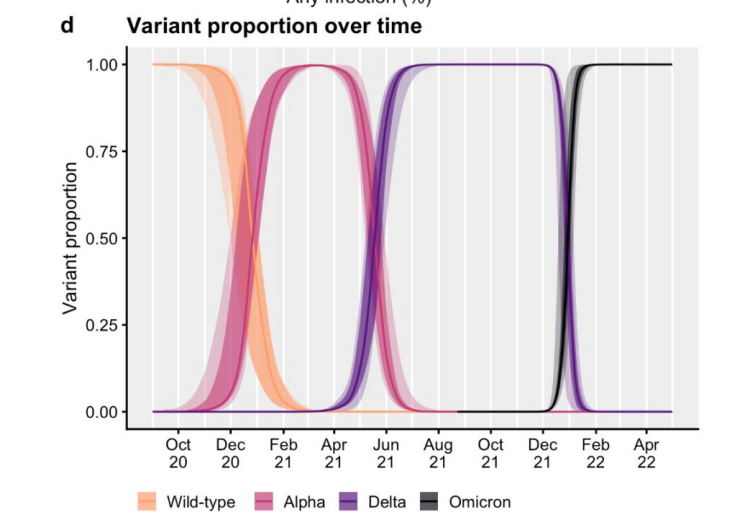
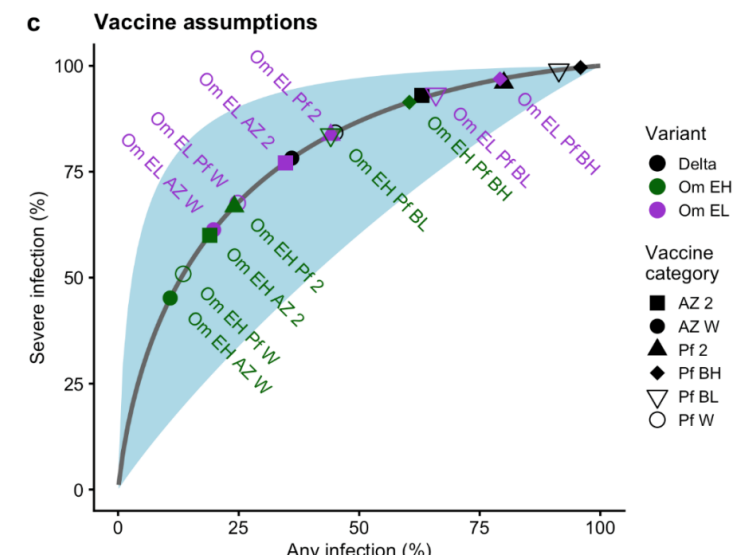


Sarbecovirus RBD ELISA (OD450)

|          | SARS-CoV-1 | RaTG13 | Pangolin-GD | Pangolin-GX | SARS-CoV-1 PC4-127 | SARS-CoV-1 Sin852 | WIV1  | Clade 3 BM48-31 | Clade 2 YN2013 |
|----------|------------|--------|-------------|-------------|--------------------|-------------------|-------|-----------------|----------------|
| Clade 1b | 3.462      | 3.447  | 3.417       | 3.405       | 3.467              | 3.476             | 0.353 | 0.135           | 0.086          |
| Clade 1a | 0.353      | 0.135  | 0.086       | 0.193       | 0.061              | 3.258             | 1.132 | 2.657           | 0.421          |
| Clade 3  | 0.996      | 0.812  | 1.95        | 0.709       | 2.889              | 3.403             | 3.598 | 3.523           | 3.473          |
| Clade 2  | 3.458      | 3.495  | 3.468       | 3.464       | 3.413              | 2.984             | 2.409 | 1.976           | 3.336          |
| Clade 3  | 0.133      | 3.424  | 0.115       | 3.483       | 3.515              | 3.476             | 0.168 | 0.244           | 0.074          |
| Clade 2  | 0.168      | 0.244  | 0.074       | 0.601       | 0.072              | 2.683             |       |                 |                |

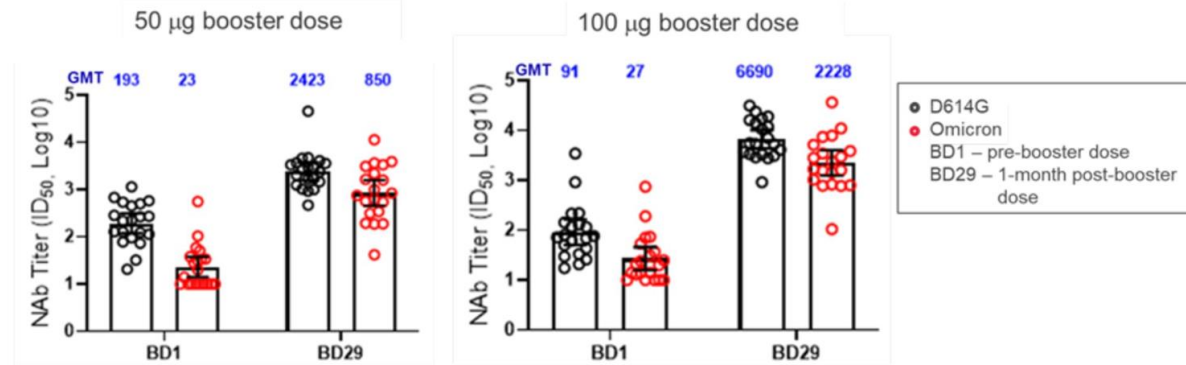


asson strategies for Omicron, due to altered receptor binding domain, coupled with enhanced stability.  
<https://www.medrxiv.org/content/10.1101/2021.12.16.472934v1.article-info>



Edmunds, Davies et. al: We use the relationship between mean neutralisation titre and protective efficacy from Khoury et al. (7) to arrive at assumptions for vaccine efficacy against infection with Omicron, given each drop in neutralisation. We then use Khoury et al.'s modelled relationship between efficacy against any infection and efficacy against severe infection to generate vaccine effectiveness estimates against severe outcomes  
[https://cmmid.github.io/topics/covid19/reports/omicron\\_england/report\\_11\\_dec\\_2021.pdf](https://cmmid.github.io/topics/covid19/reports/omicron_england/report_11_dec_2021.pdf)

## Sera from recipients of 100 µg primary series who were boosted with 50 or 100 µg (20/dose)

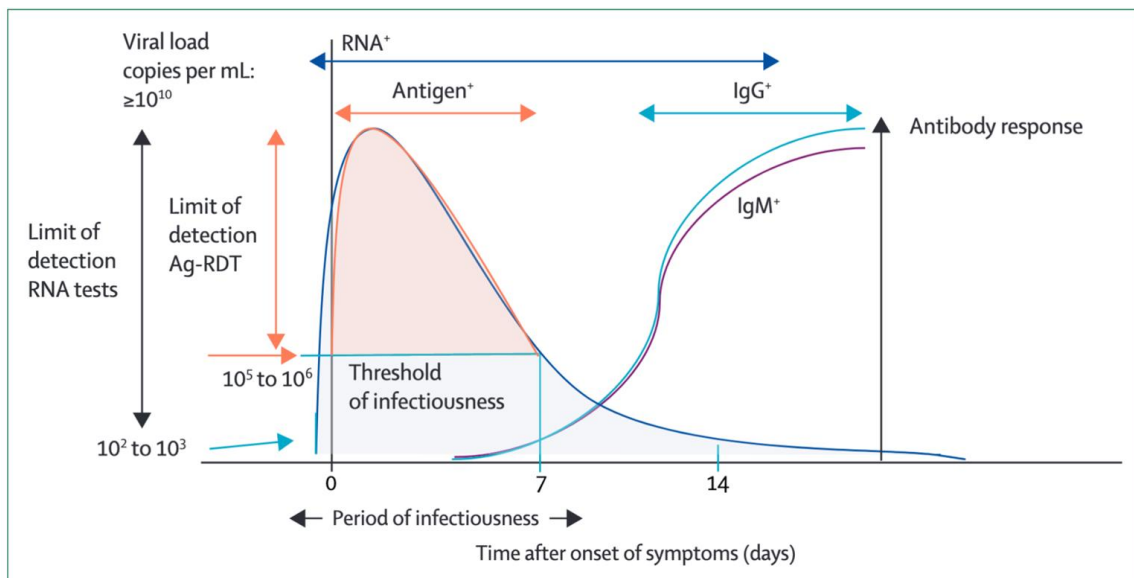


- Neutralizing antibody titers to Omicron are increased 4 weeks post 50 µg or 100 µg booster dose of mRNA-1273 (~37 & ~83-fold increases, respectively)

Moderna Data on File

Moderna press update yesterday

<https://twitter.com/erictopol/status/1472869635250814980>

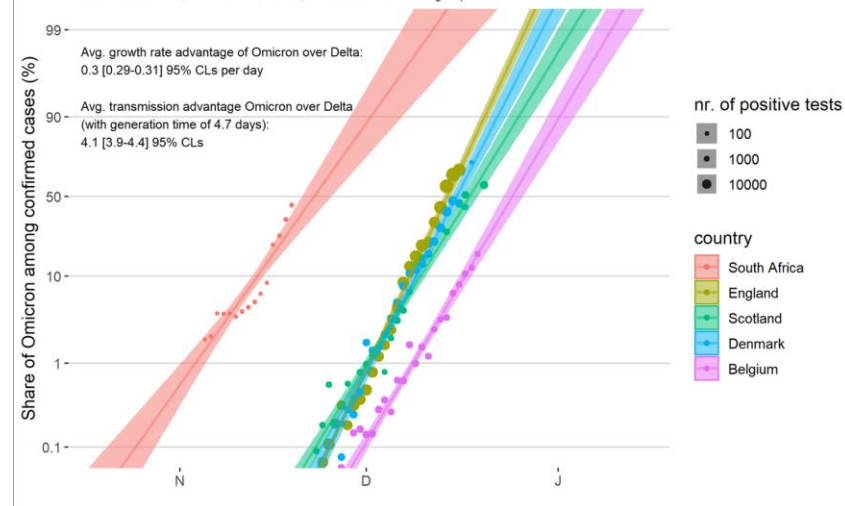


Evidence from 113 studies done in 17 countries shows that SARS-CoV-2 viral RNA can be detected as early as 6 days before symptom onset, concentrations peak around the time of symptom onset or a few days later, and it usually becomes undetectable from upper respiratory tract samples about 2 weeks after symptom onset, and with no substantial differences between adults and children.<sup>17</sup> The viral load from lower respiratory tract samples might be higher, peak later, and persist for longer than the load from upper respiratory tract samples.

<https://www.thelancet.com/action/showPdf?pii=S0140-6736%2821%2902346-1>

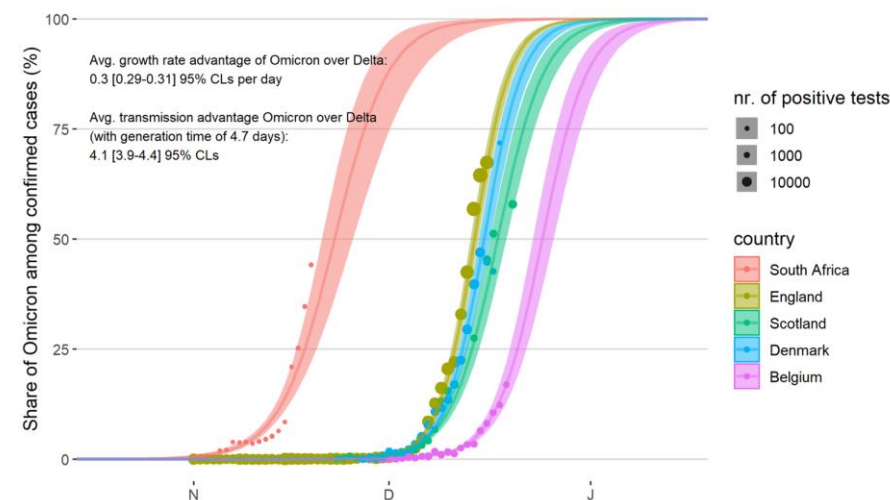
## Spread of Omicron variant inferred from S gene target failure (SGTF) data (SA, UK, BE) & variant PCR data (DK)

(SGTF counts adjusted by proportion that was estimated to be Omicron based on GISAID data, data Lesley Scott & NHLS team, UKHSA, Public Health Scotland, Statens Serum Institut, Federal Test Platform Belgium)



## Spread of Omicron variant inferred from S gene target failure (SGTF) data (SA, UK, BE) & variant PCR data (DK)

(SGTF counts adjusted by proportion that was estimated to be Omicron based on GISAID data, data Lesley Scott & NHLS team, UKHSA, Public Health Scotland, Statens Serum Institut, Federal Test Platform Belgium)

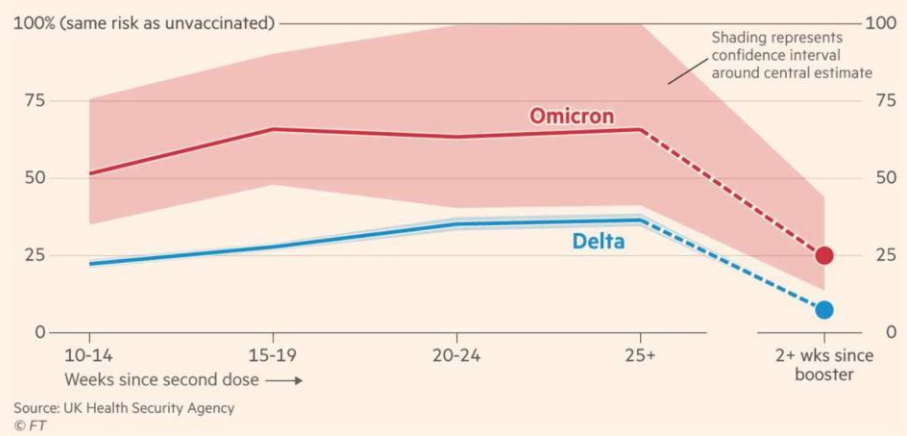


<https://twitter.com/twenseleers/status/1472946174973915145>



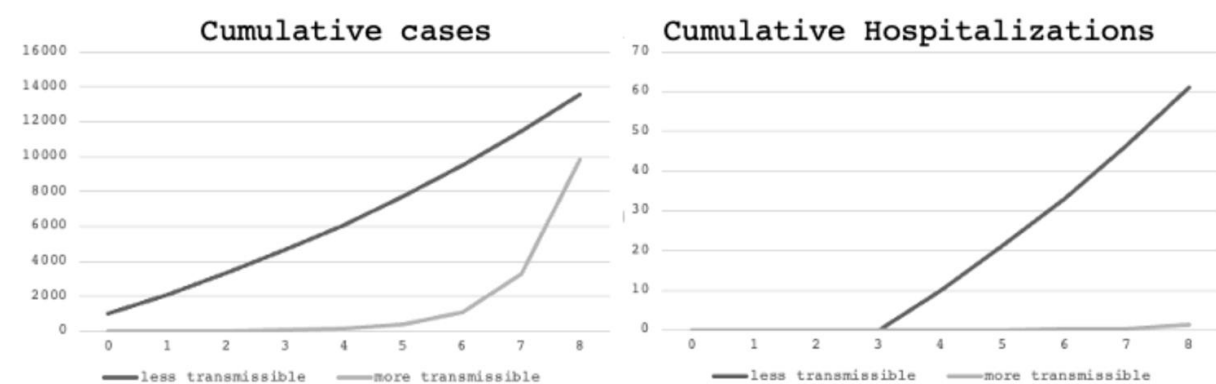
English data shows substantially increased risk of breakthrough infections with Omicron, but boosters push risk back down towards Delta levels

Relative risk of symptomatic infection for someone with the Pfizer/BioNTech vaccine compared with an unvaccinated person



Potential to expect 5x more breakthrough cases after booster Omicron to Delta.

<https://twitter.com/erlichya/status/1470759683233546244?s=21>



<https://twitter.com/billhanage/status/1470613668404604935?s=21>

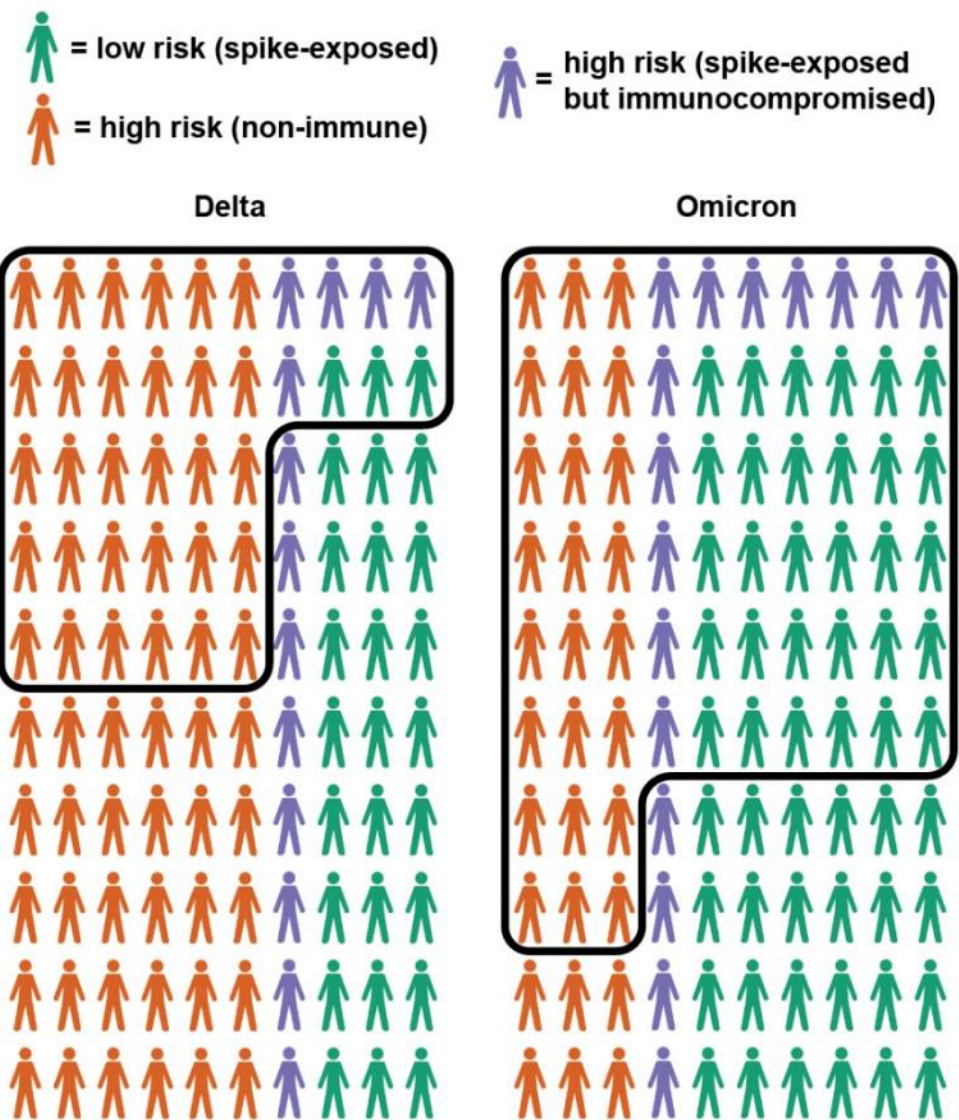


Figure 1. Challenges in comparing IFR of Omicron vs Delta

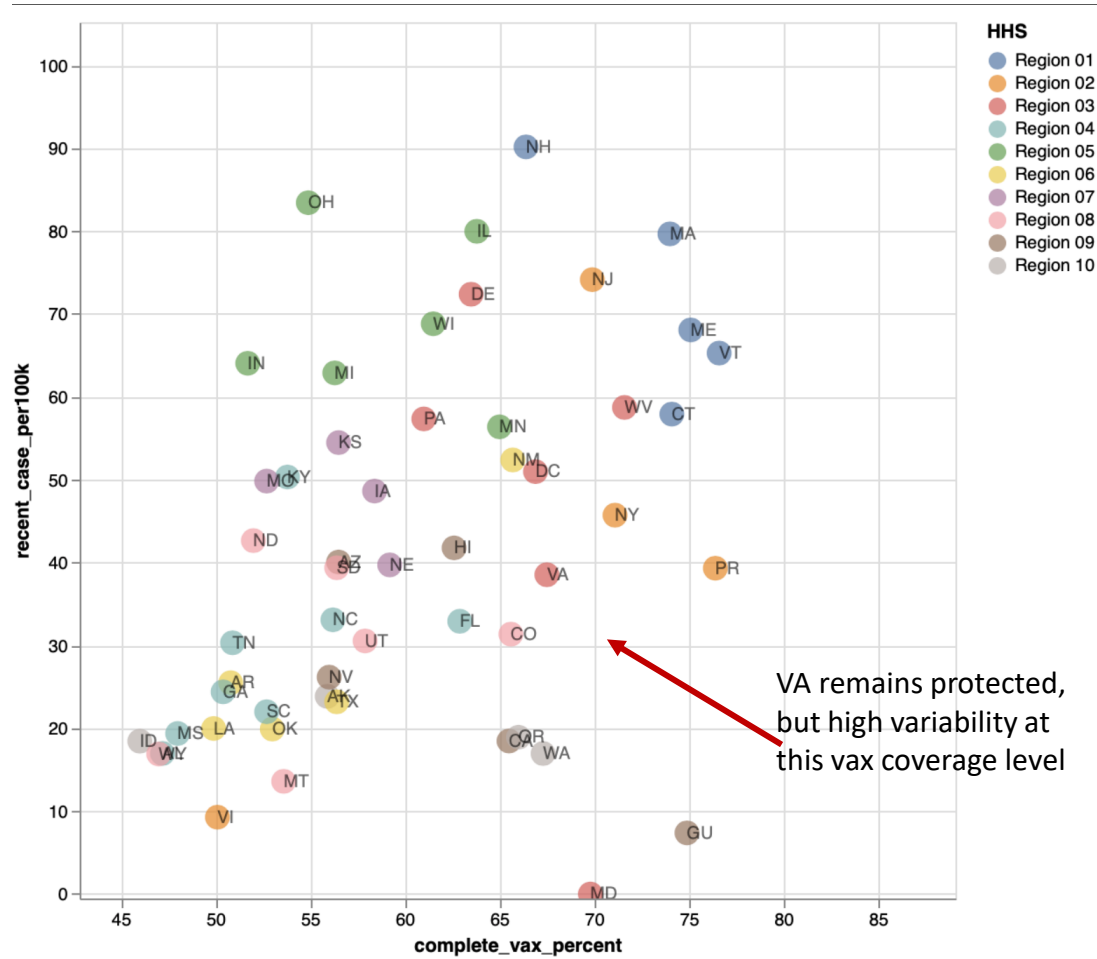
Harvard study highlights bias challenges in estimating severity

[https://twitter.com/roby\\_bhatt/status/1471212023576870912](https://twitter.com/roby_bhatt/status/1471212023576870912)

# Recent Cases Correlate with Vax Coverage

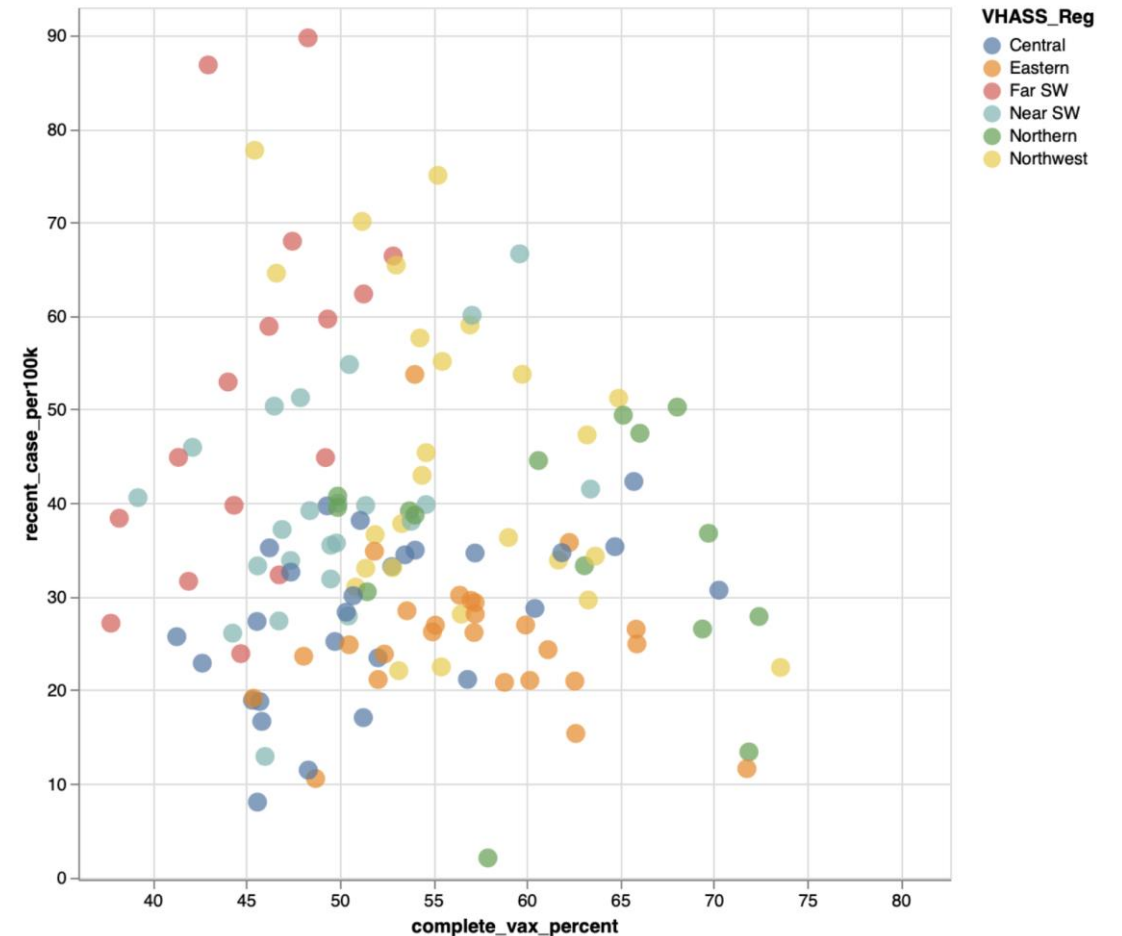
## Mean cases per 100K vs. vaccine coverage

- Correlations between vax coverage and recent case rates has disappeared as more high coverage states have high rates



## Virginia Counties

- Counties with higher vax coverage slightly lower rates

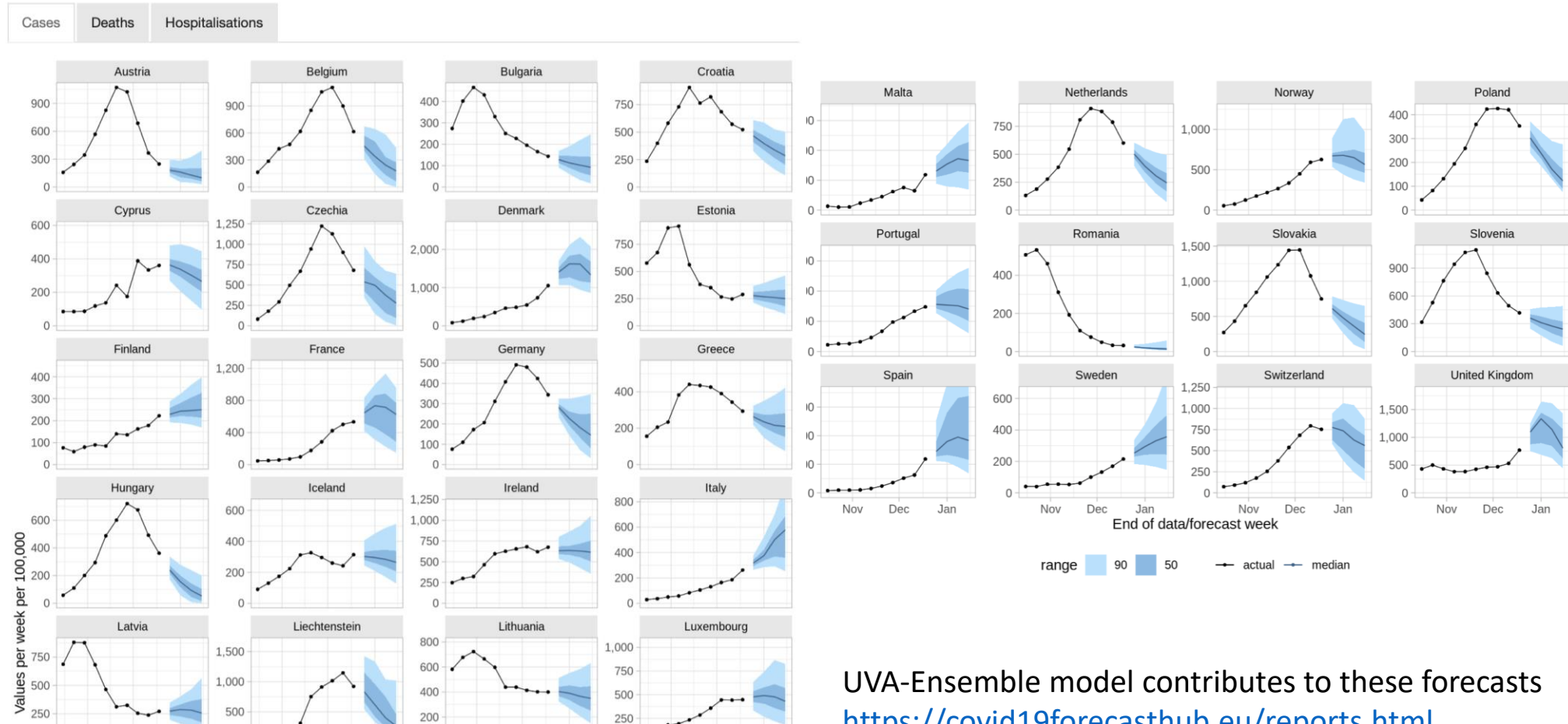


# European Nations

European COVID-19 Forecast Hub Evaluation Report  
EuroCOVIDhub-ensemble

## Latest forecasts

Forecasts of cases/deaths per week per 100,000. Click the **Forecast** tab above to view all past forecasts.



- Case Rate changes are mixed in Europe with many countries growing rapidly while other decline
- Growth and high rates in most nations reporting significant prevalence of Omicron

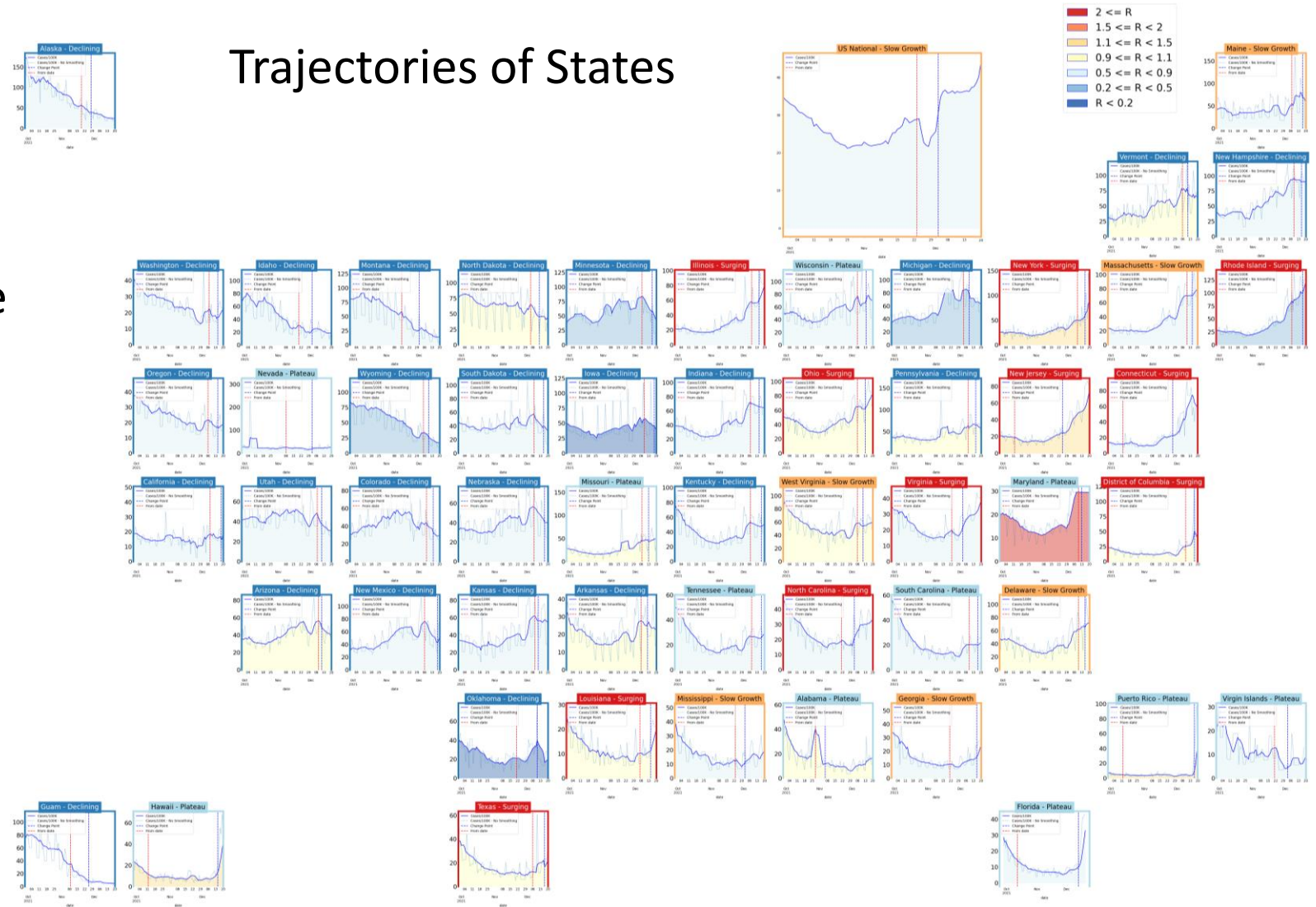
UVA-Ensemble model contributes to these forecasts  
<https://covid19forecasthub.eu/reports.html>



# United States Overall

- Overall growth has paused
- Significant number of states remain in growth
- Case rates remain moderate to high in most states

## Trajectories of States

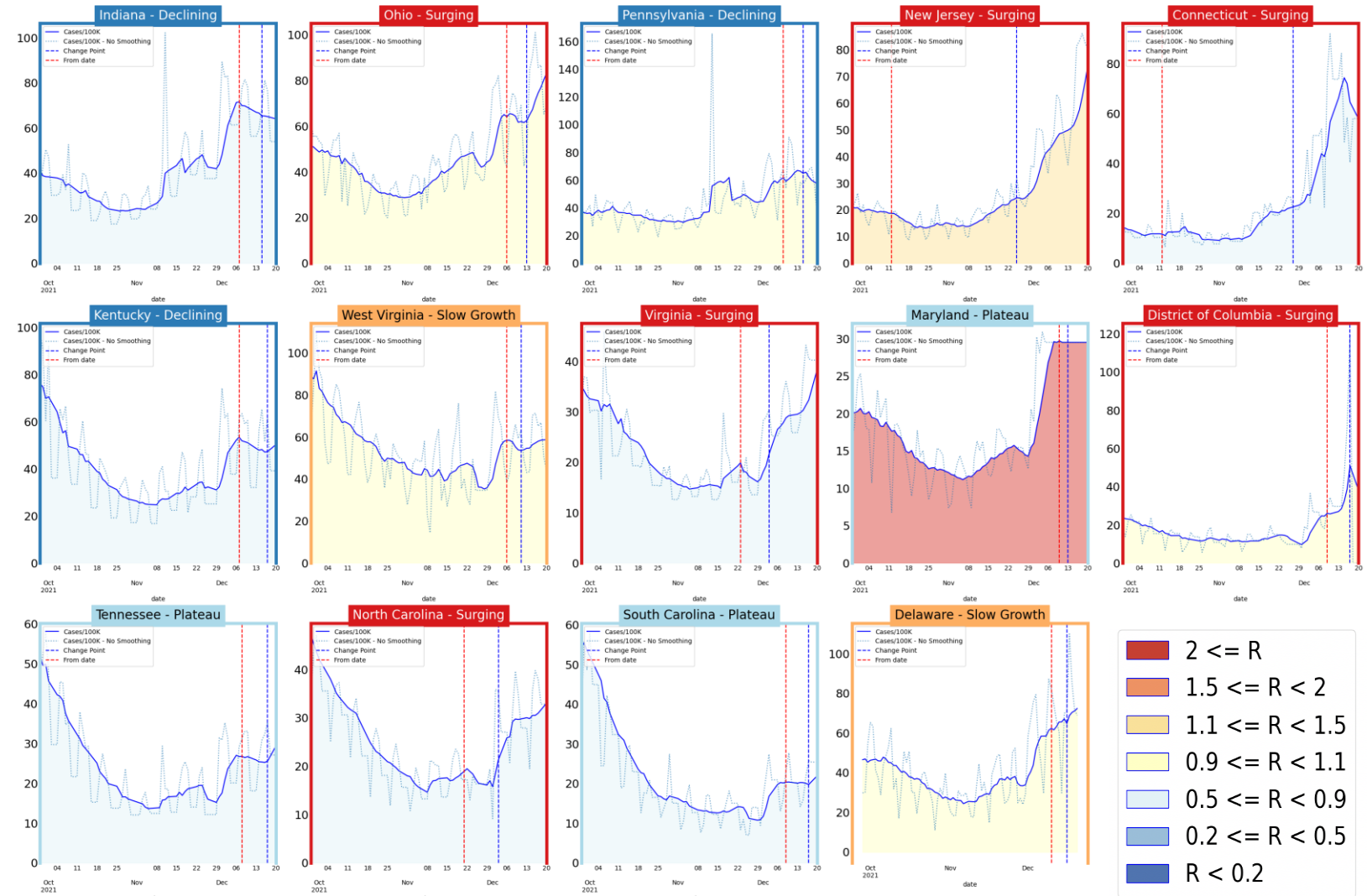


| Status      | # States (1 week ago) |
|-------------|-----------------------|
| Declining   | 26 (22)               |
| Plateau     | 11 (13)               |
| Slow Growth | 6 (5)                 |
| In Surge    | 11 (14)               |



# Virginia and Her Neighbors

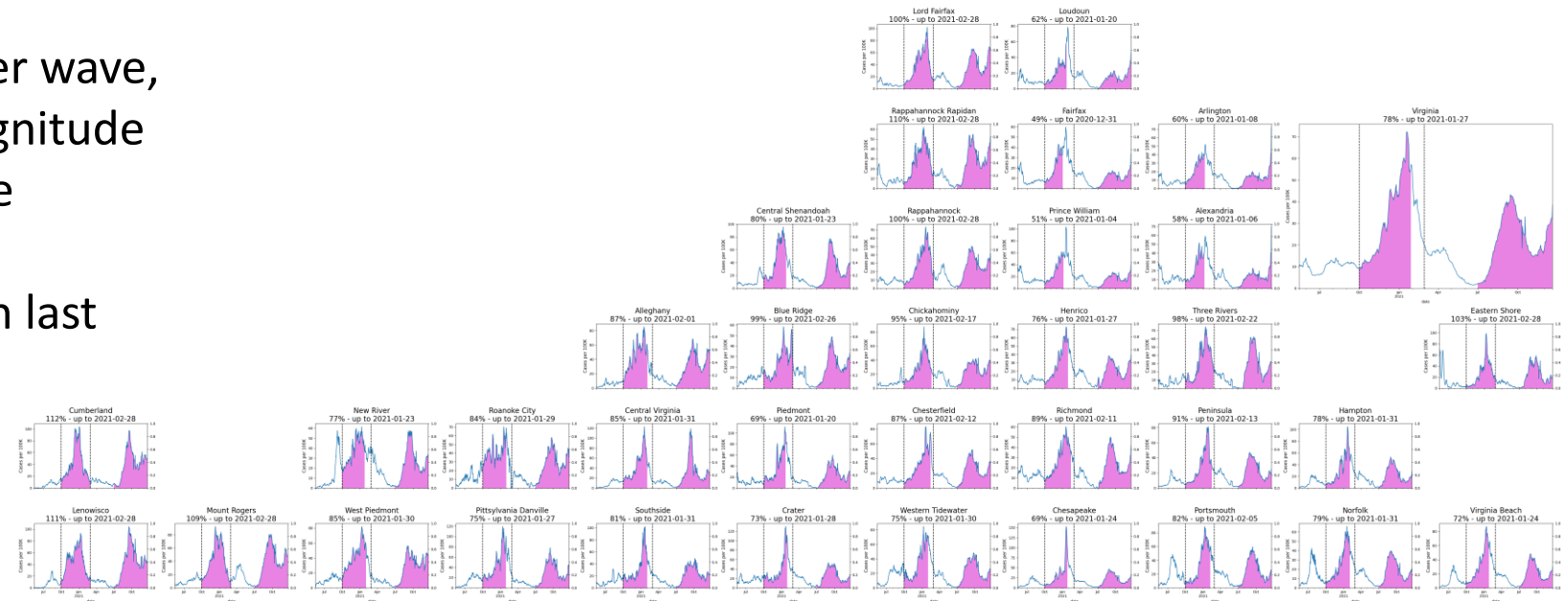
- Recent case rate growth seems have paused in VA and most neighbors
- Case rates remain high as arrival of Omicron may fuel more growth



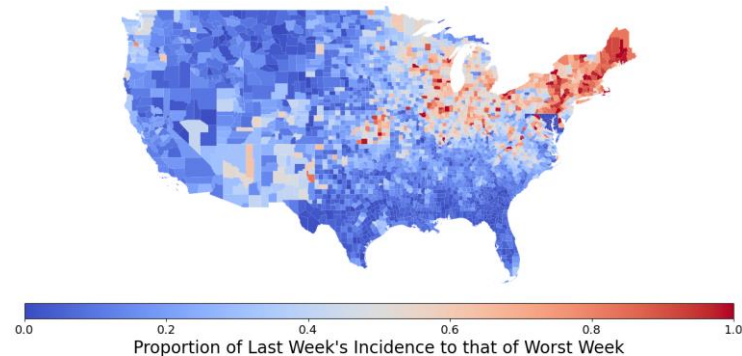
# Virginia and Her Neighbors

- Current Delta and Fall- Winter wave, quickly approach overall magnitude of last years Fall-Winter wave
- Many districts have already experienced more cases than last year's Fall-Winter

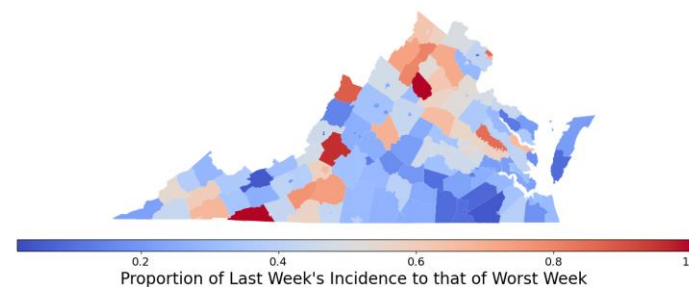
Matched Period of Current Cumulative Case Rate to Previous Surge



Recent Incidence Compared to Worst Week by County



Recent Incidence Compared to Worst Week by County



- Some Counties approaching and experiencing the worst week of the pandemic this last week
- Nationally worst weeks are concentrated in Northeast

# Zip code level weekly Case Rate (per 100K)

## Case Rates in the last week by zip code

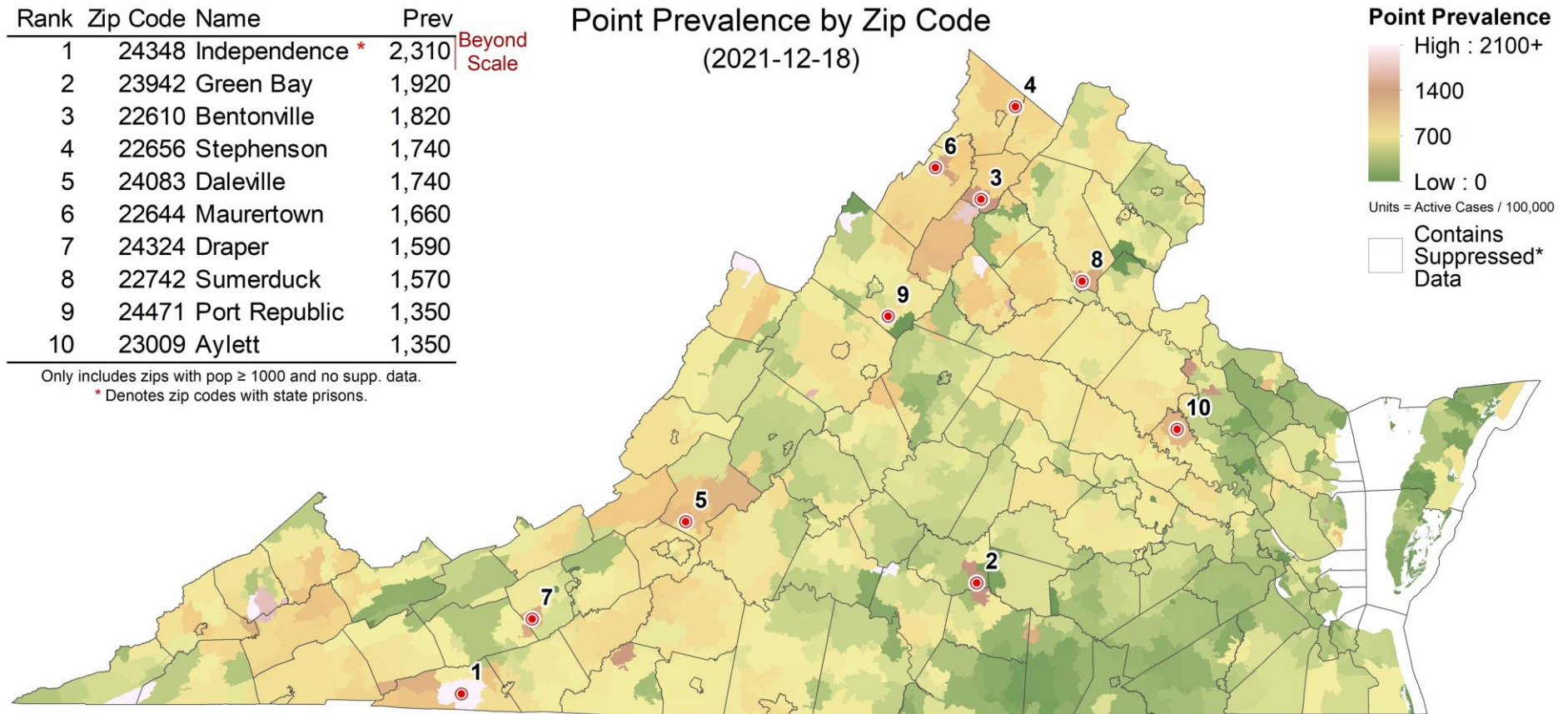
- Color scaled adjusted to accommodate the very high prevalence levels this week
- Clusters of high prevalence in Southwest and Northwest
- Some counts are low and suppressed to protect anonymity, those are shown in white

| Rank | Zip Code | Name           | Prev  |
|------|----------|----------------|-------|
| 1    | 24348    | Independence * | 2,310 |
| 2    | 23942    | Green Bay      | 1,920 |
| 3    | 22610    | Bentonville    | 1,820 |
| 4    | 22656    | Stephenson     | 1,740 |
| 5    | 24083    | Daleville      | 1,740 |
| 6    | 22644    | Maurertown     | 1,660 |
| 7    | 24324    | Draper         | 1,590 |
| 8    | 22742    | Sumerduck      | 1,570 |
| 9    | 24471    | Port Republic  | 1,350 |
| 10   | 23009    | Aylett         | 1,350 |

Only includes zips with pop ≥ 1000 and no supp. data.

\* Denotes zip codes with state prisons.

Point Prevalence by Zip Code  
(2021-12-18)



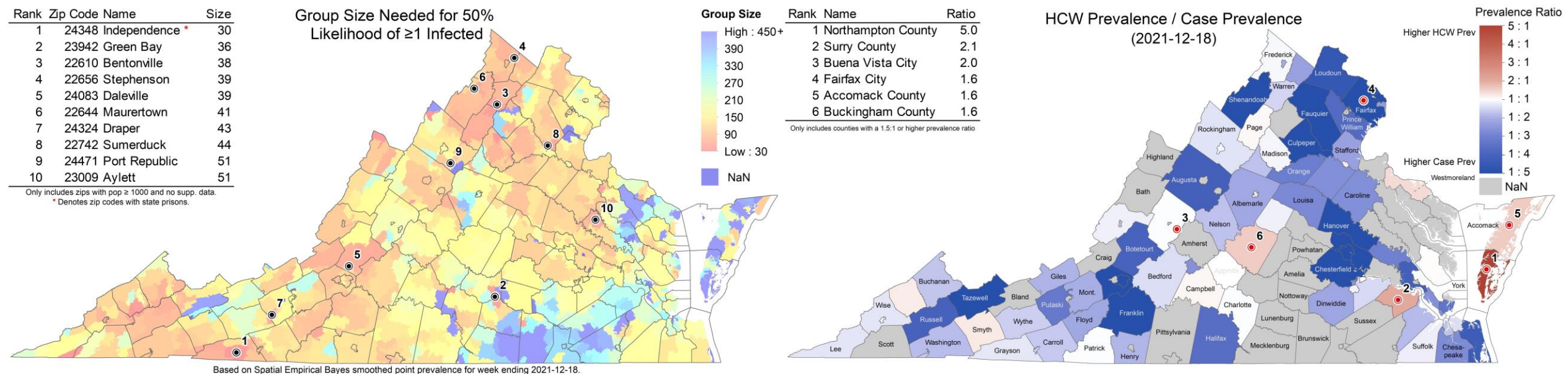
Based on Spatial Empirical Bayes smoothed point prevalence for week ending 2021-12-18.



# Risk of Exposure by Group Size and HCW prevalence

## Case Prevalence in the last week by zip code used to calculate risk of encountering someone infected in a gathering of randomly selected people (group size 25)

- **Group Size:** Assumes 2 undetected infections per confirmed case (ascertainment rate from recent seroprevalence survey), and shows minimum size of a group with a 50% chance an individual is infected by zip code (eg in a group of 30 in Stephenson, there is a 50% chance someone will be infected)
- **HCW ratio:** Case rate among health care workers (HCW) in the last week using patient facing health care workers as the denominator / general population's case prevalence

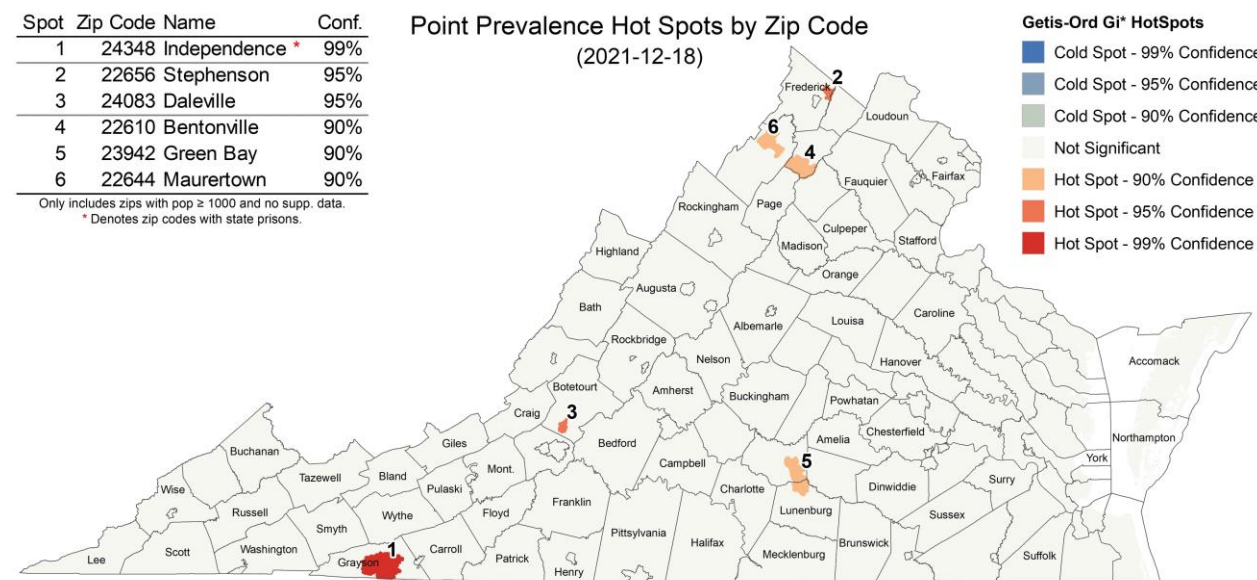


# Current Hot-Spots

## Case rates that are significantly different from neighboring areas or model projections

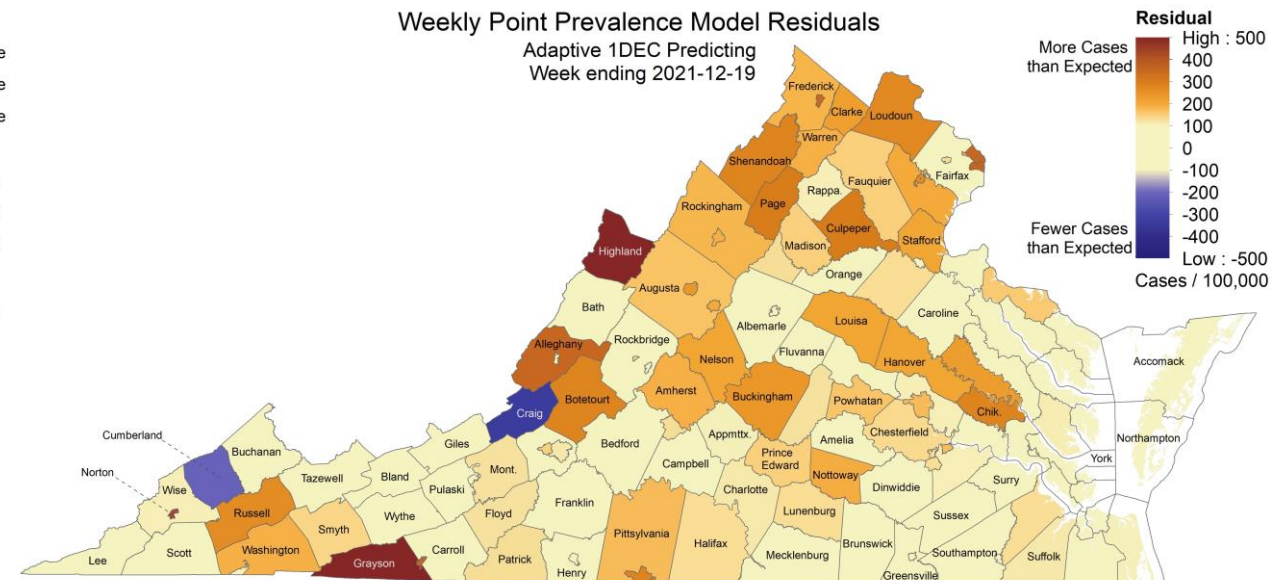
- **Spatial:** Getis-Ord Gi\* based hot spots compare clusters of zip codes with weekly case prevalence higher than nearby zip codes to identify larger areas with statistically significant deviations
- **Temporal:** The weekly case rate (per 100K) projected last week compared to observed by county, which highlights temporal fluctuations that differ from the model's projections

### Spatial Hotspots



Based on Global Empirical Bayes smoothed point prevalence for week ending 2021-12-18.

### Clustered Temporal Hotspots



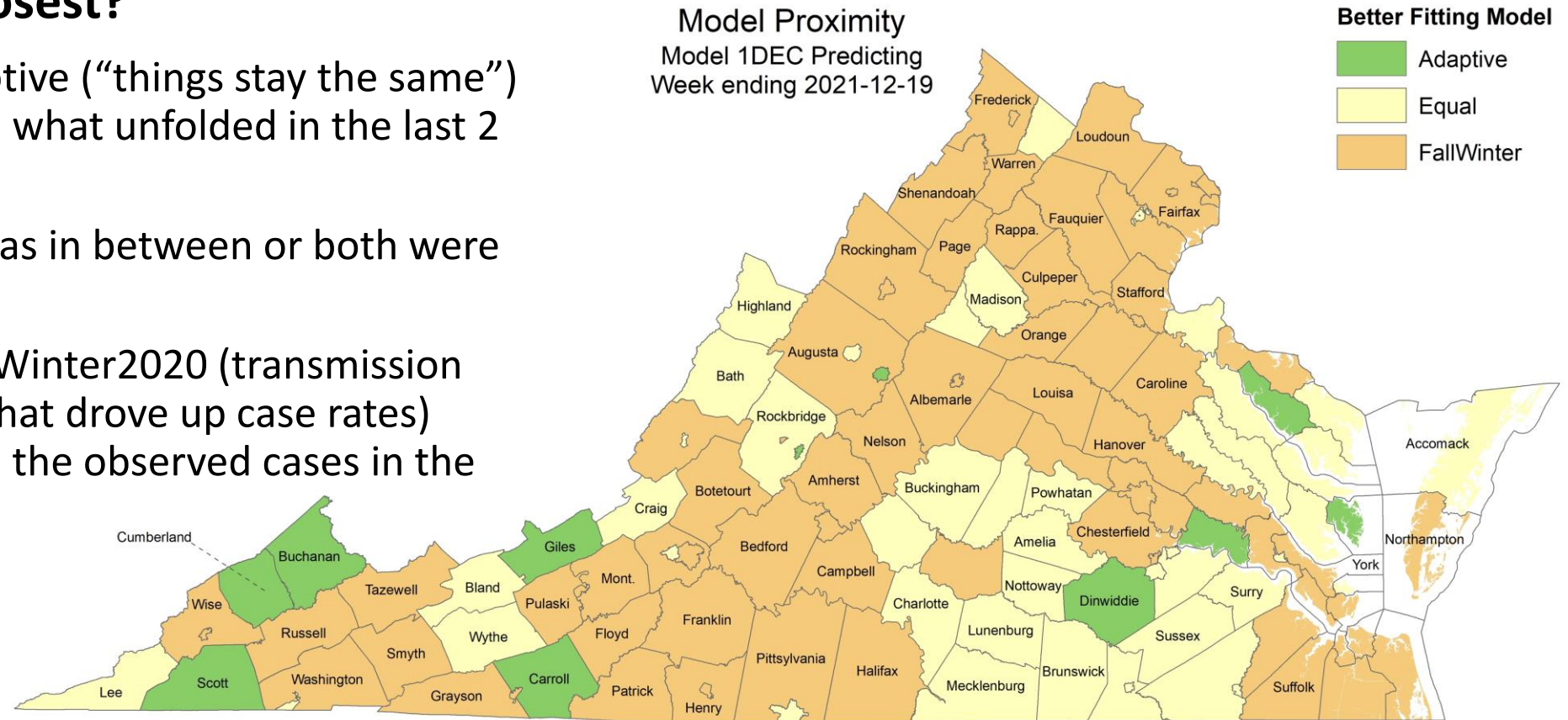
Moran's I = 0.027028, Z-Score = 1.530613, P-Value = 0.125865  
No Residual Autocorrelation Detected



# Scenario Trajectory Tracking

## Which scenario from last projection did each county track closest?

- Green means the Adaptive (“things stay the same”) scenario was closest to what unfolded in the last 2 weeks
- Yellow means reality was in between or both were very similar
- Orange means the FallWinter2020 (transmission drivers from last year that drove up case rates) scenario was closest to the observed cases in the last 2 weeks



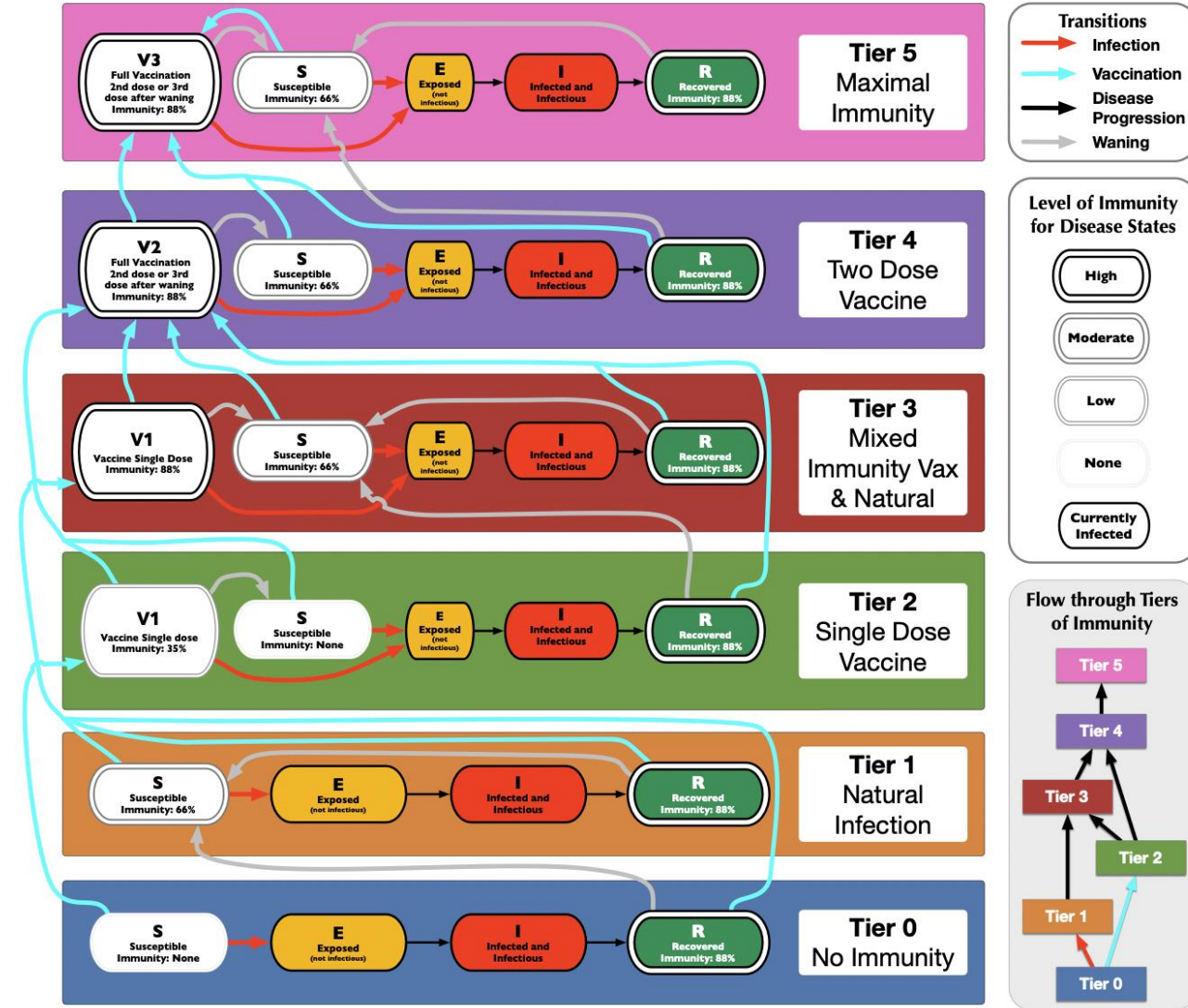
# Model Update – Adaptive Fitting

---

# New Model Structure Focused on Tiers of Immunity

## Uncertainty surrounds the rate of waning immunity

- New model structure built to better track levels and timing of waning
- Outcomes vary based on age and immune history; for partial immunity, protection against hospitalization and death is stronger than No Immunity but weaker than Maximal Immunity
- Use same Adaptive fitting approach with vaccine schedules and simulated infections driving movement across the tiers
- Different Scenarios can also be applied





# Adaptive Fitting Approach

## Each county fit precisely, with recent trends used for future projection

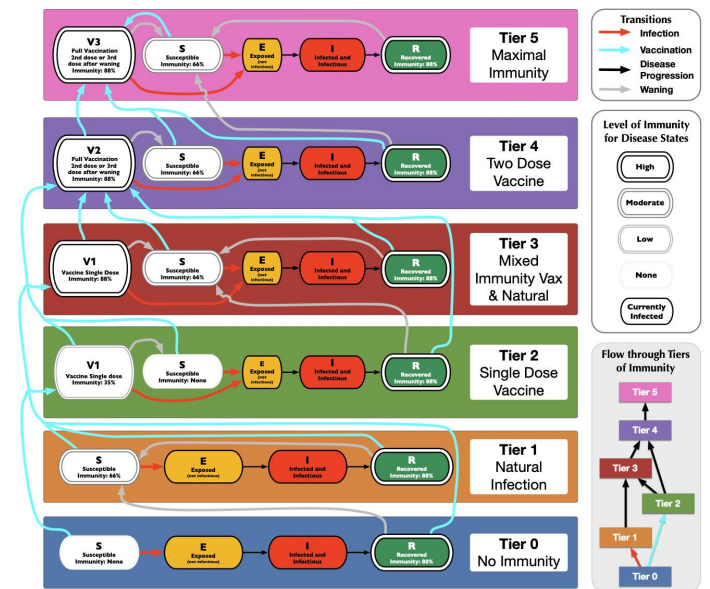
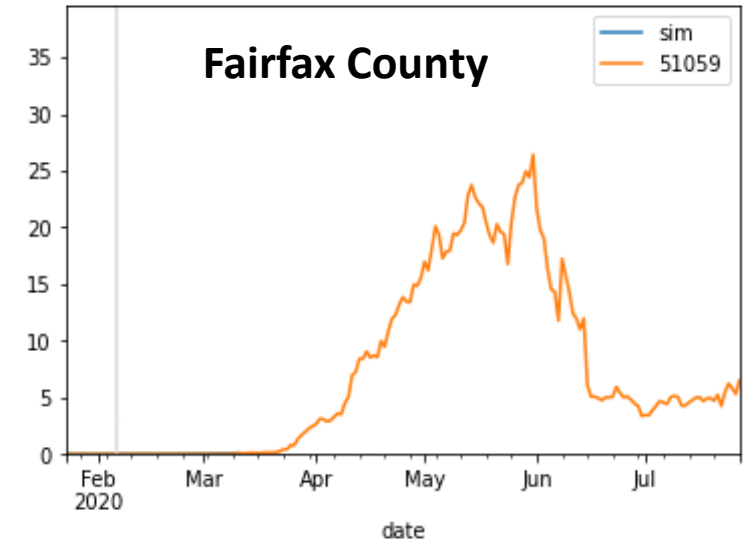
- Allows history to be precisely captured, and used to guide bounds on projections

## Model: An alternative use of the same meta-population model, PatchSim with multiple tiers of immunity

- Allows for future “what-if” Scenarios to be layered on top of calibrated model
- Allows for waning of immunity and for partial immunity against different outcomes (eg lower protection for infection than death)

## External Seeding: Steady low-level importation

- Widespread pandemic eliminates sensitivity to initial conditions, we use steady 1 case per 10M population per day external seeding



# Using Ensemble Model to Guide Projections

Ensemble methodology that combines the Adaptive v machine learning and statistical models such as:

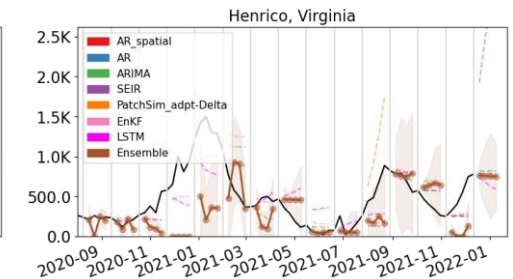
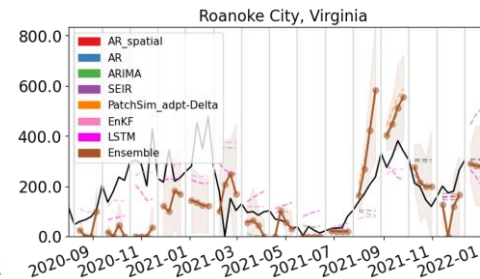
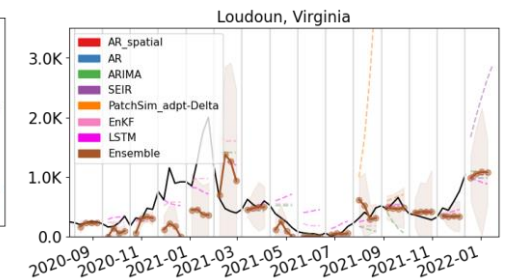
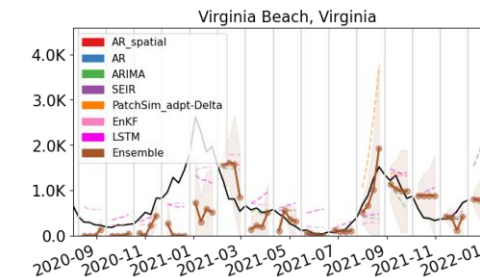
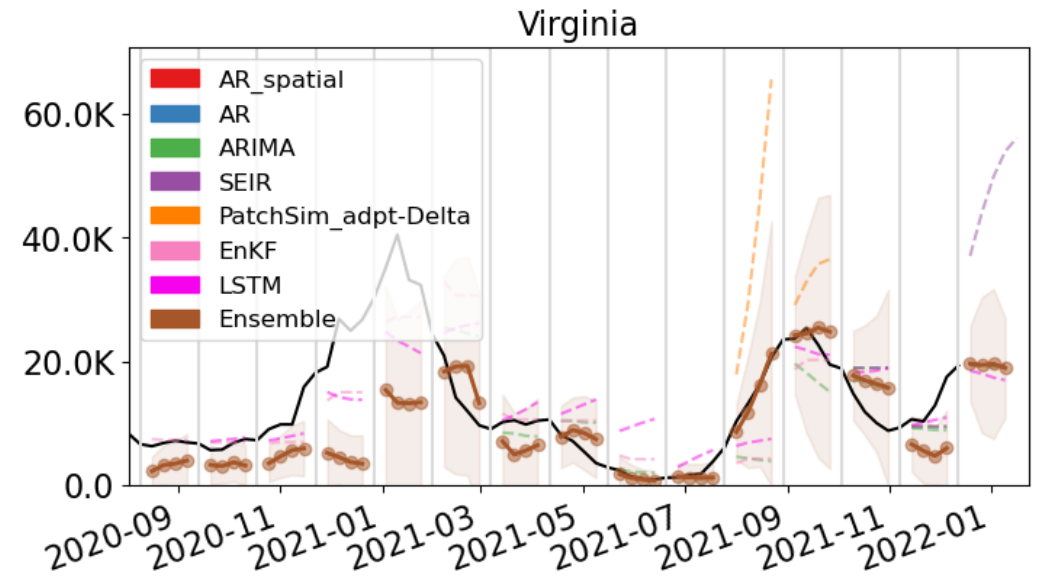
- Autoregressive (AR, ARIMA)
- Neural networks (LSTM)
- Kalman filtering (EnKF)

Weekly forecasts done at co

Models chosen because track record in disease forecasting and to in diversity and robustness.

Ensemble fore provides additional 'surveillance' for making sce

Also s



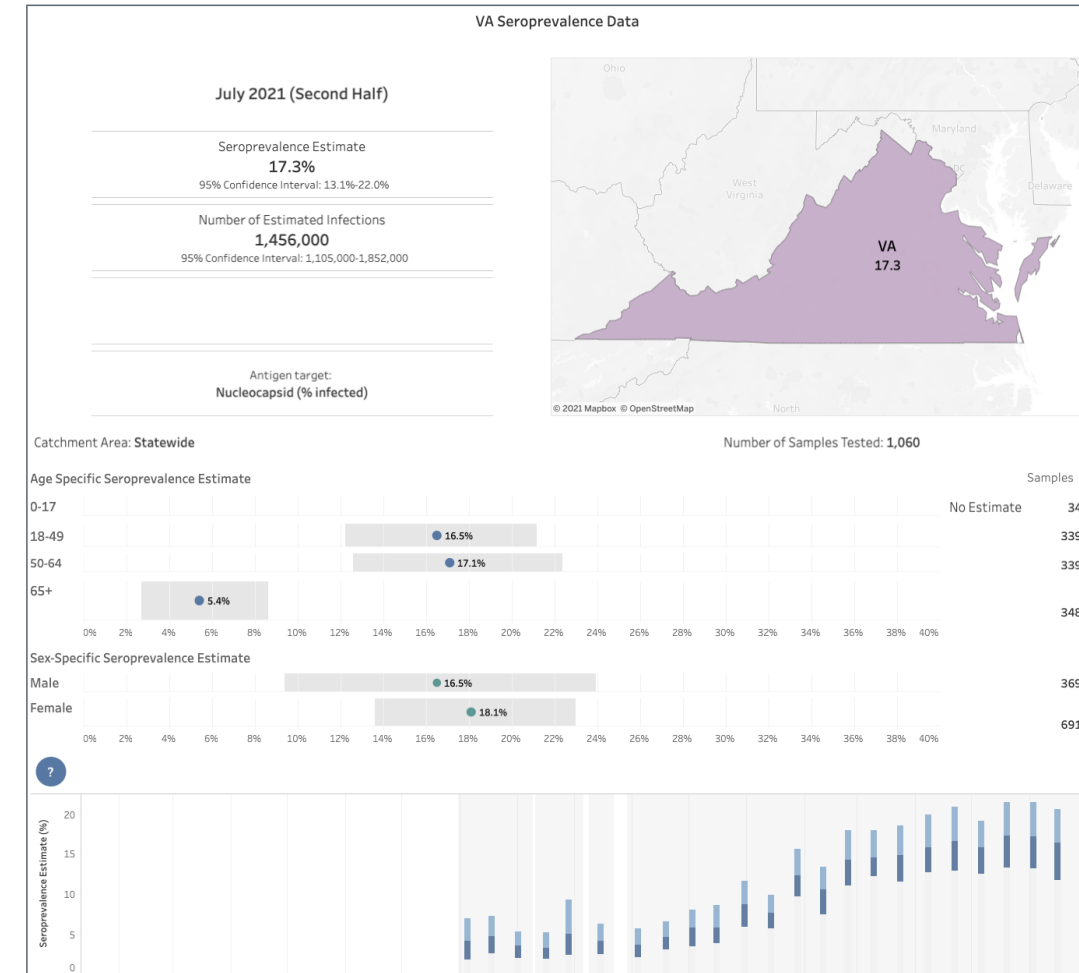
# Seroprevalence updates to model design

**Several seroprevalence studies provide better picture of how many actual infections have occurred**

- CDC Nationwide Commercial Laboratory Seroprevalence Survey

**These findings are equivalent to an ascertainment ratio of ~2x in the future, with bounds of (1.3x to 3x)**


- Thus for 2x there are 2 total infections in the population for every confirmed case recently
- **Case ascertainment is half of that for those with prior immunity**
- Uncertainty design has been shifted to these bounds (previously higher ascertainment as was consistent earlier in the pandemic were being used)



<https://covid.cdc.gov/covid-data-tracker/#national-lab>

# Calibration Approach

- **Data:**
  - County level case counts by date of onset (from VDH)
  - Confirmed cases for model fitting
- **Calibration:** fit model to observed data and ensemble's forecast
  - Tune transmissibility across ranges of:
    - Duration of incubation (5-9 days), infectiousness (3-7 days)
    - Undocumented case rate (1x to 7x) guided by seroprevalence studies
    - Detection delay: exposure to confirmation (4-12 days)
  - Approach captures uncertainty, but allows model to precisely track the full trajectory of the outbreak
- **Project:** future cases and outcomes generated using the collection of fit models run into the future
  - **Mean trend from last 7 days of observed cases and first week of ensemble's forecast used**
  - Outliers removed based on variances in the previous 3 weeks
  - 2 week interpolation to smooth transitions in rapidly changing trajectories
- **Outcomes:** Data driven by shift and ratio that has least error in last month of observations
  - Hospitalizations: 3 days from confirmation, 6.8% of cases hospitalized
  - Deaths: 11 days from confirmation, 1.45% of cases die



# COVID-19 in Virginia:

Dashboard Updated: 12/14/2021

Data entered by 5:00 PM the prior day.

Cases, Hospitalizations and Deaths

Total Cases\*

1,003,110

(New Cases: 2,416)^

Confirmed†

736,568

Probable†

266,542

Total Hospitalizations\*\*

40,234

Confirmed†

37,828

Probable†

2,406

Total Deaths

14,992

Confirmed†

12,578

Probable†

2,414

\* Includes people with either a positive molecular/PCR test (Confirmed), positive antigen test (Probable) or symptomatic with known exposure to COVID-19 (Probable).

\*\* Hospitalization of a case is captured at the time VDH performs case investigation. This underrepresents the total number of hospitalizations in Virginia.

^New cases represent the number of confirmed and probable cases reported to VDH in the past 24 hours.

† VDH adopted the updated CDC COVID-19 2021 Surveillance Case Definition on September 1, 2021 which is found here: -- <https://ndc.services.cdc.gov/case-definitions/coronavirus-disease-2019-2021/>

Source: Cases - Virginia Electronic Disease Surveillance System (VEDSS), data entered by 5:00 PM the prior day.

Outbreaks

Total Outbreaks\*

5,786

Outbreak Associated Cases

94,723

\* At least two (2) lab confirmed cases are required to classify an outbreak.

Testing (PCR Only)

Testing Encounters PCR Only\*

10,775,815

Current 7-Day Positivity Rate PCR Only\*\*

8.7%

\* PCR refers to "Reverse transcriptase polymerase chain reaction laboratory testing."

\*\* Lab reports may not have been received yet. Percent positivity is not calculated for days with incomplete data.

Multisystem Inflammatory Syndrome in Children

Total Cases\*

123

Total Deaths

1

\*Cases defined by CDC HAN case definition: <https://emergency.cdc.gov/han/2020/han00432.asp>

Accessed 8:30am December 15, 2021  
<https://www.vdh.virginia.gov/coronavirus/>

# Scenarios – Transmission Conditions

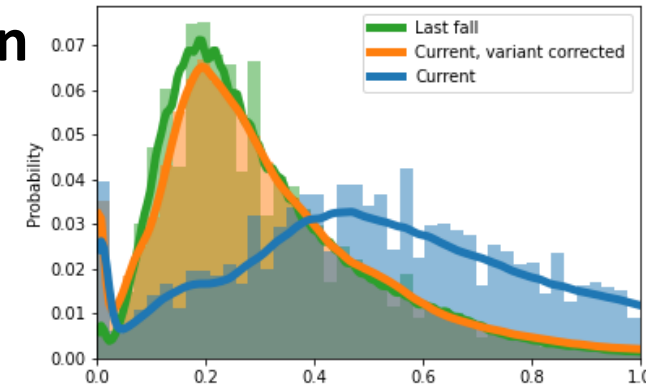
- Variety of factors continue to drive transmission rates
  - Seasonal impact of weather patterns, travel and gatherings, fatigue and premature relaxation of infection control practices
- **Waning Immunity:** Mean of 6 months to a re year protection (rate of 0.0027) similar to [Pfizer study](#)
- **Projection Scenarios:**
  - **Adaptive:** Control remains as is currently experienced into the future with assumption that Delta remains as the majority strain
  - **Adaptive-Omicron:** Controls remain the same while the Omicron rapidly dominates prevalence. Has same transmissibility as Delta with 30% immune evasion
  - **Adaptive-FallWinter:** Starting this week the core drivers of transmission from Sept 2020 – Feb 2021 are coarsely replayed but boosted to account for Delta's increased transmissibility
  - **Adaptive-Surge Control:** Starting in one week behaviors and mitigation efforts ramp up over a 2-week period culminating in a 25% reduction in transmission



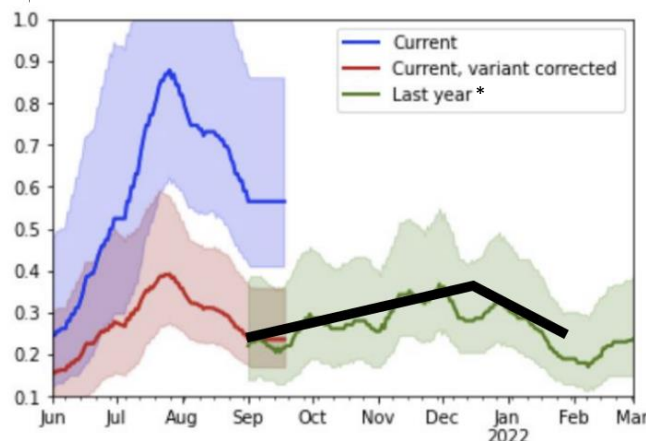
# Scenarios – FallWinter Description

## September 2020 – February 2021 saw a strong wave of transmission

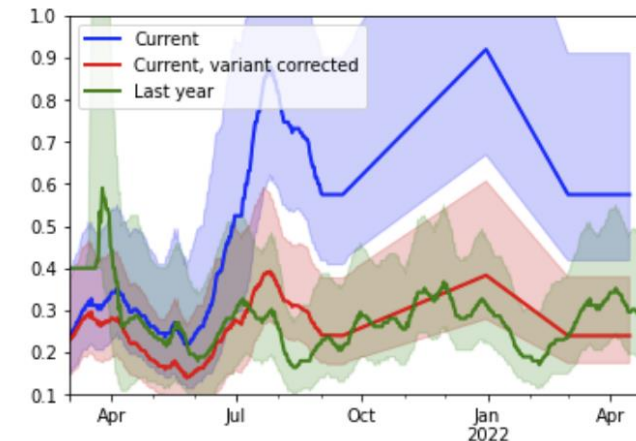
- We analyze previous Fall-Winter's wave vs. current Delta driven wave and observe surprising similarities
  - The distribution of fitted model transmissibility is nearly identical between these periods when corrected for Delta's increased transmissibility
- **FallWinter** tries to capture the “transmission drivers” from the past and use them as if they were to occur again this season but with Delta variant (compared to ancestral)
  - Use the above analysis of fitted model transmissibilities from Sept 2020 – Feb 2021 to guide the future transmissibility from Sept 2021 through Feb 2022, but add the enhanced transmissibility of Delta back in



**Fitting:**  
Black line  
represents the  
coarsely fitted  
base  
transmissibility



\* “Last year” is transplanted into 2021-22



**Delta enhanced:**  
Blue trajectory  
represents current  
fitted and then  
projected  
transmissibility in  
FallWinter2020

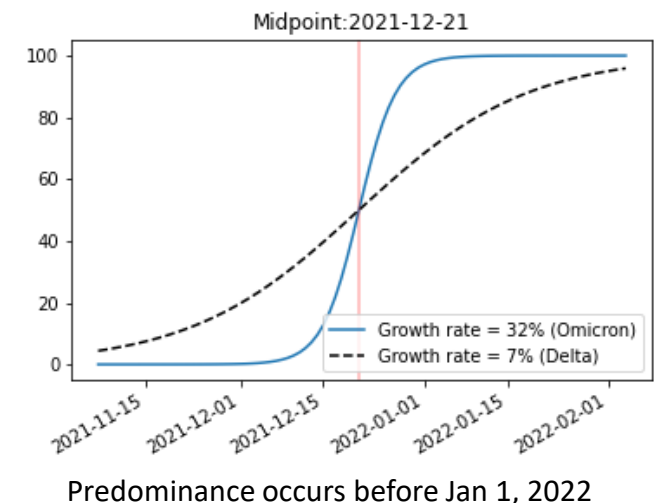
# Scenarios – Omicron Description

## Omicron shown ability to evade immunity and may be more transmissible

- **Transmissibility:** Evidence exists that it may be more inherently transmissible, for this scenario we conservatively keep Omicron with the **same transmissibility of Delta**
- **Immune Evasion:** Stronger evidence demonstrates that Omicron can cause infection in those with some immunity (natural and vaccine induced). Conservative estimate of **30% immune evasion** allows Omicron to infect 30% of individuals that would have otherwise been protected against Delta
- **Prevalence:** Proportion of cases caused by Omicron variant estimated from growth rates observed in other countries with similar levels of immunity (growth of 32%, doubling in ~3 days)
- **Severity:** Initial reports suggest Omicron may not cause as severe disease as Delta, we use a 50% reduction in severity for hospitalizations and deaths.
  - Recent [preprint from South Africa](#) suggests it could be 70% less, whereas, a [PHE report](#) found no evidence of reduced severity.

**These are conservative estimates (lower) for both transmissibility and immune evasion, even so growth is stronger than previously observed**

## Estimated Prevalence curve for US



# Scenarios – Vaccination Conditions

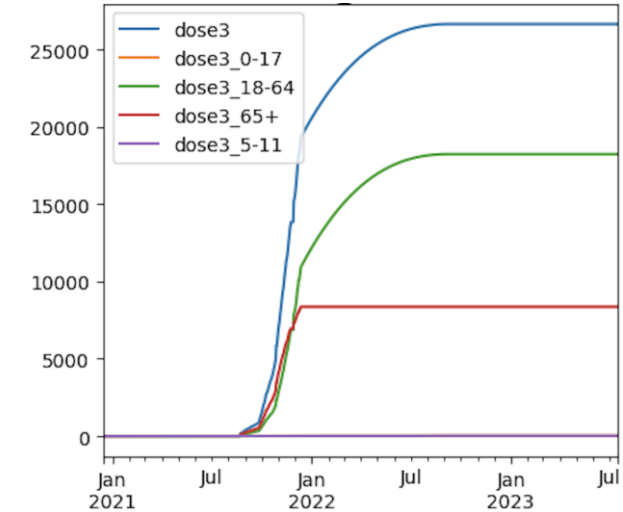
## Vaccine Characteristics

- **Pfizer/Moderna:** 50% after first dose, 95% after second dose (3.5 week gap) **J & J :** 67% efficacy after first dose
- Delay to efficacy from doses is 14 days, immunity lasts at least 7m ([NEJM study](#))

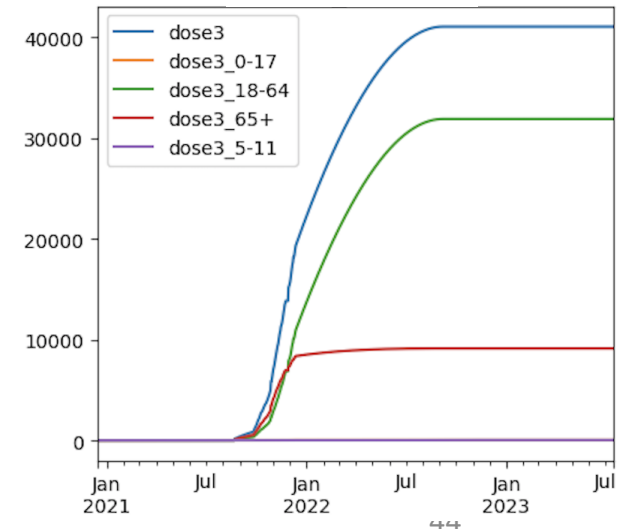
## Vaccine Administration Scenarios

- **Status quo (no label):**
  - **Eventual coverage:** COVIDcast corrected acceptance estimates (statewide mean is ~80% adults, 65% of population) reached by end of January.
  - **Children (5-11):** Follow rates of 12-16 year olds, max out at 80% of adult acceptance
  - **3<sup>rd</sup> doses:** Top out with total coverage of 40%
- **Optimistically High Boosting (HighBoost):**
  - **Eventual coverage:** COVIDcast corrected acceptance estimates (statewide mean is ~80% adults, 65% of population) reached by end of January.
  - **Children (5-11):** Follow rates of 12-16 year olds, max out at 80% of adult acceptance
  - **3<sup>rd</sup> doses:** Top out with total coverage of 70%
- Acceptance at county level = regional acceptance +/- relative current vax
- Front-loaded rollout (two-thirds of the remaining in half the time)

Status Quo



High Boost



# Projection Scenarios – Combined Conditions

| Name                  | Txm Controls | Vax | Description  |
|-----------------------|--------------|-----|--|
| Adaptive              | C            | SQ  | Likely trajectory based on conditions remaining similar to the current experience  |
| Adaptive-HighBoost    | C            | VO  | Vaccination through January 2022 reaches an optimistically high level of expanded coverage (85%)   |
| Adaptive-Omicron      | C            | SQ  | Assumes rapid dominance of immune evading variant. Conservatively uses no transmission advantage to Delta but 30% of previously immune individuals are susceptible to infection from Omicron |
| Adaptive-SurgeControl | 25%          | SQ  | Transmission rates in the next month reduced through increased control from non-pharmaceutical interventions, with status quo vax and Delta  |
| Adaptive-FallWinter   | FallWinter   | SQ  | Transmission rates coarsely follow the rates from last September through this February but are boosted by Delta's enhanced transmissibility  |

## Transmission Controls:

C = Current levels persist into the future

25% = Transmission rates are reduced by 25% with a gradual introduction, concluding in 4 weeks

FallWinter2020 = Transmission rates from Sept 2020 – Feb 2021 are coarsely replayed but boosted by Delta's increased transmissibility

## Vaccinations:

SQ = Status quo acceptance leads to low rates of vaccination through the summer

VO = Vaccination acceptance optimistically expands with increased rates through the summer



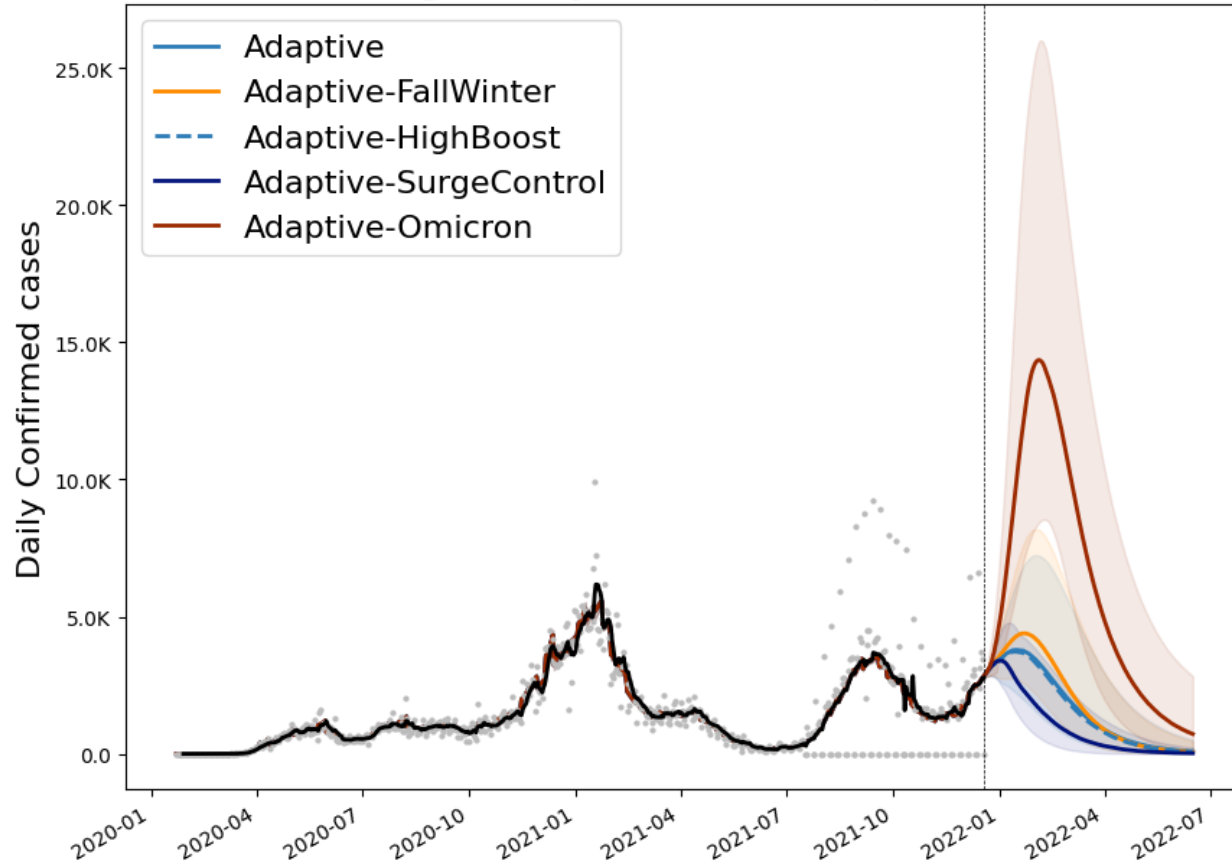
# Model Results

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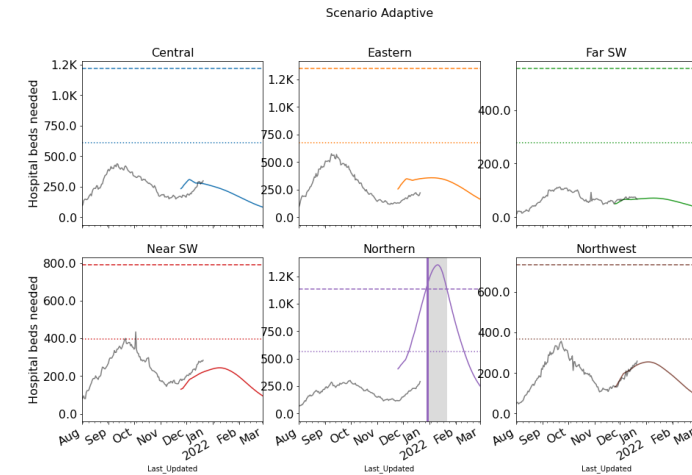
# Outcome Projections

## Confirmed cases

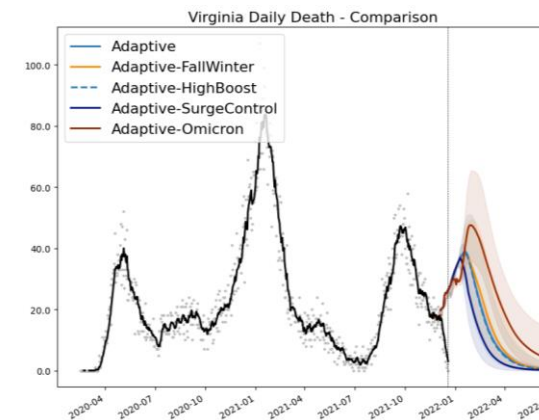
Virginia Daily Confirmed - Comparison



## Estimated Hospital Occupancy

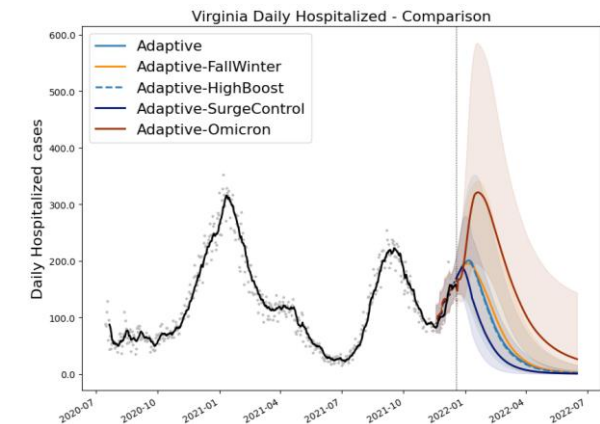


## Daily Deaths



Death ground truth from VDH "Event Date" data, most recent dates are not complete

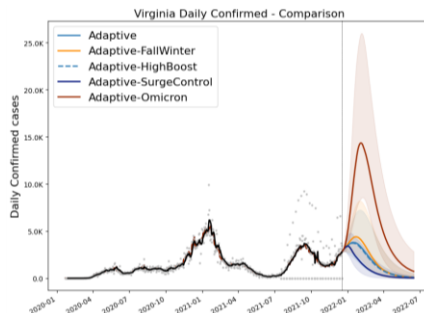
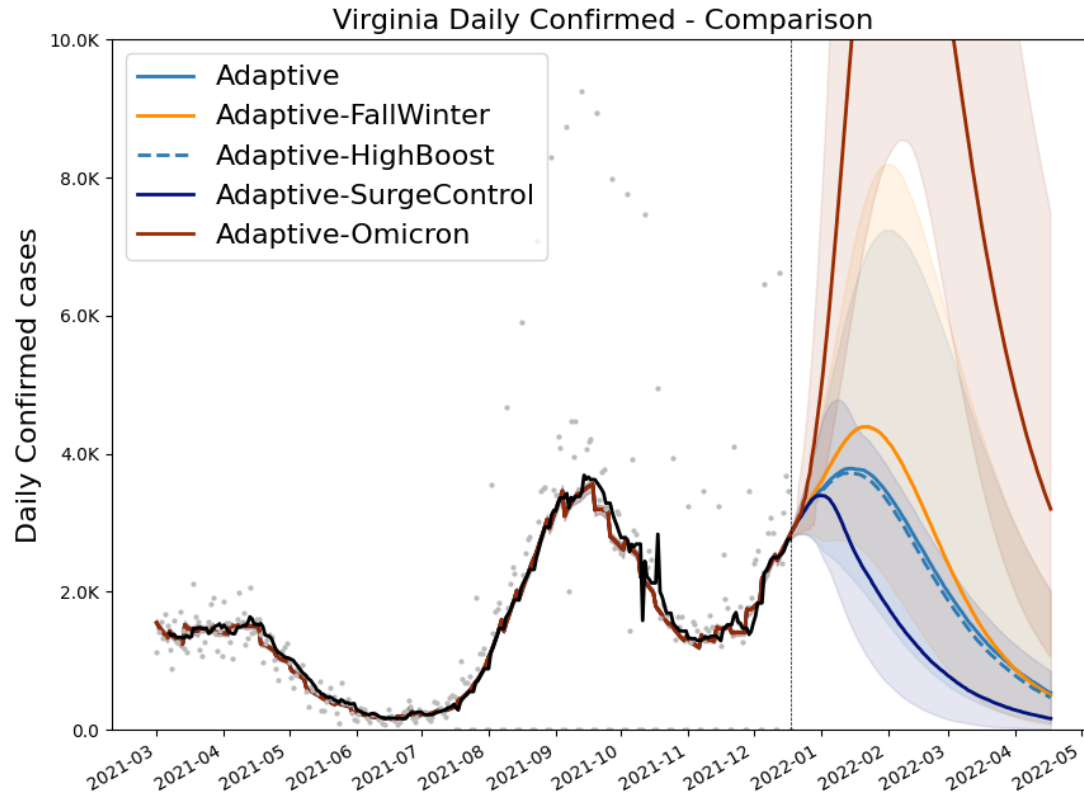
## Daily Hospitalized



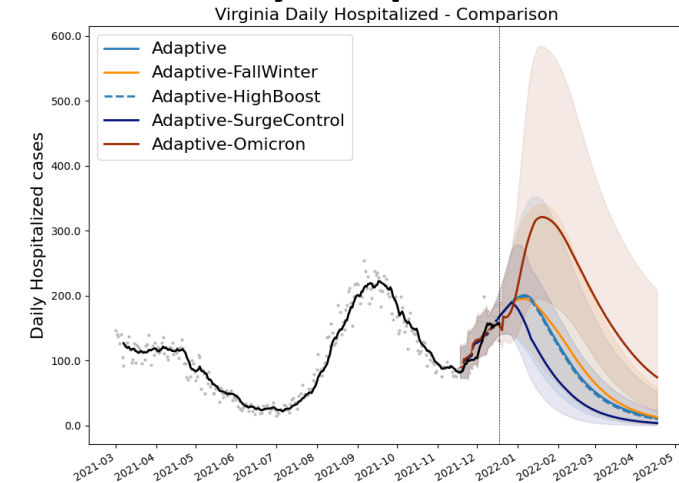
Hospitalization is slightly miscalibrated this week, seems to be driven by Northern region. Trends should be correct but absolute numbers are misaligned

# Outcome Projections – Closer Look

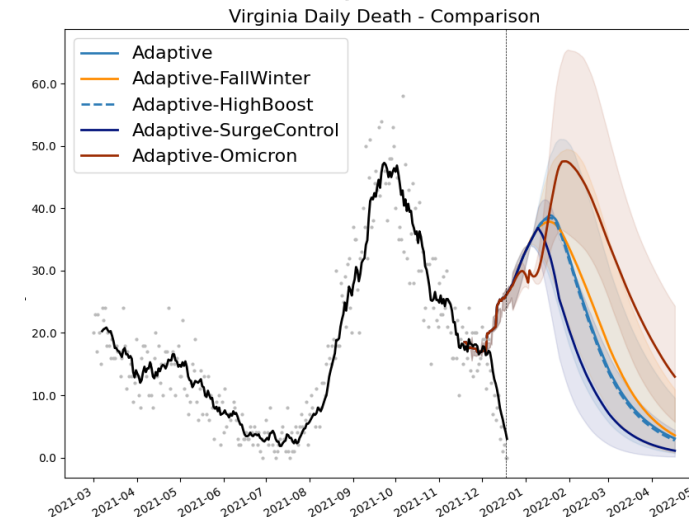
## Confirmed cases



## Daily Hospitalized



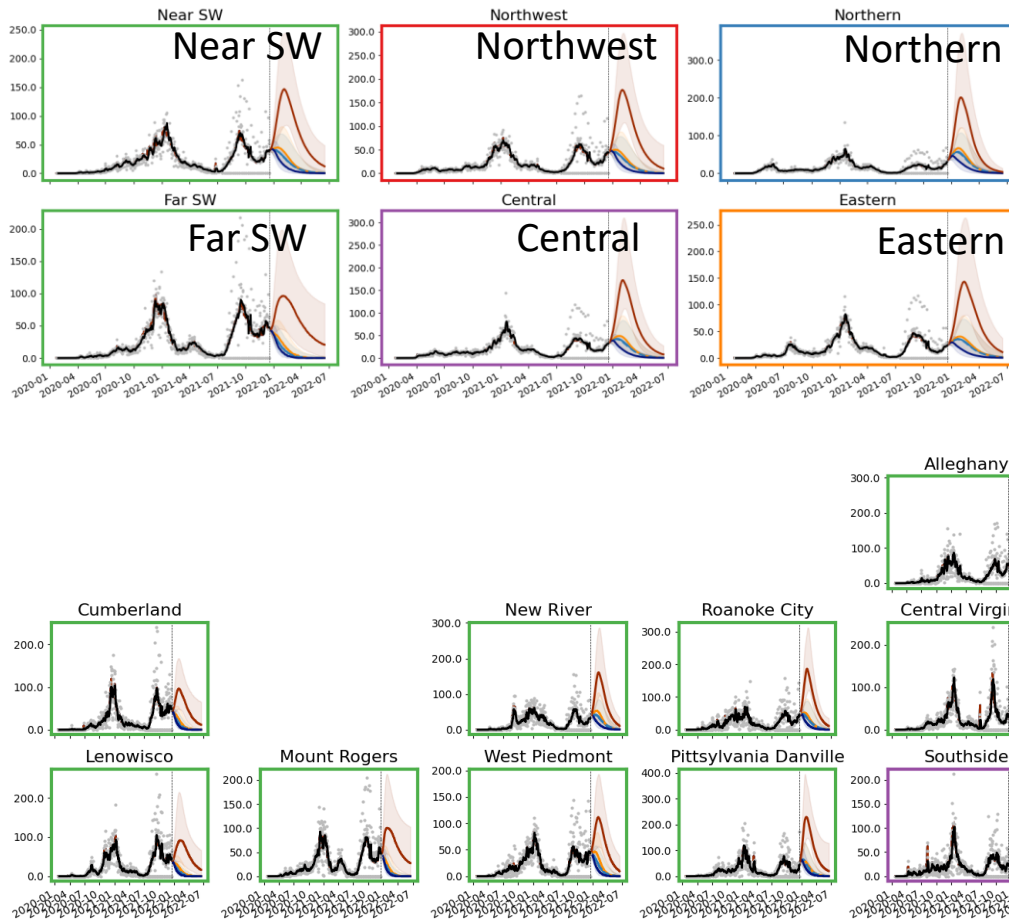
## Daily Deaths



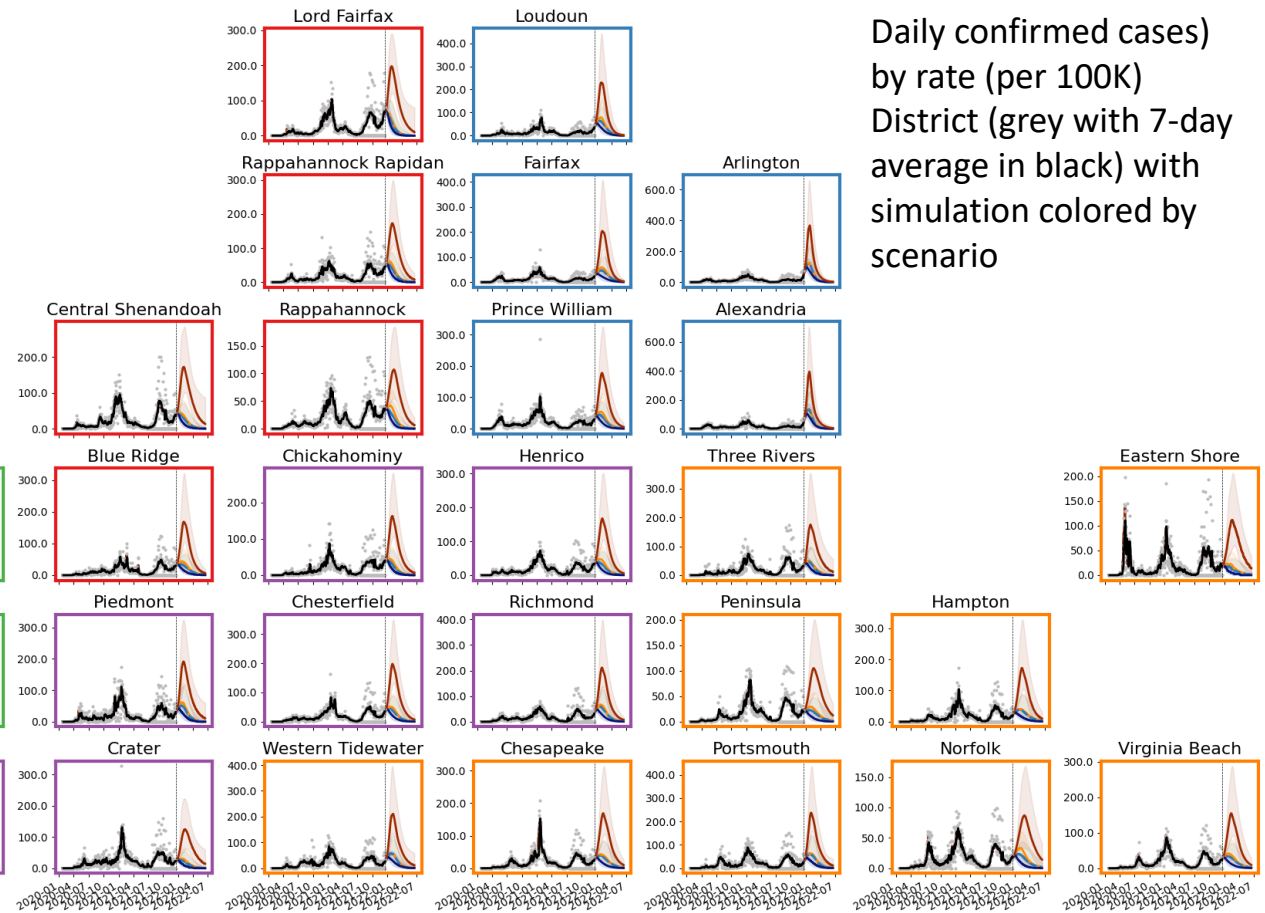
Death ground truth from VDH "Event Date"  
data, most recent dates are not complete

# Detailed Projections: All Scenarios

## Projections by Region



## Projections by District

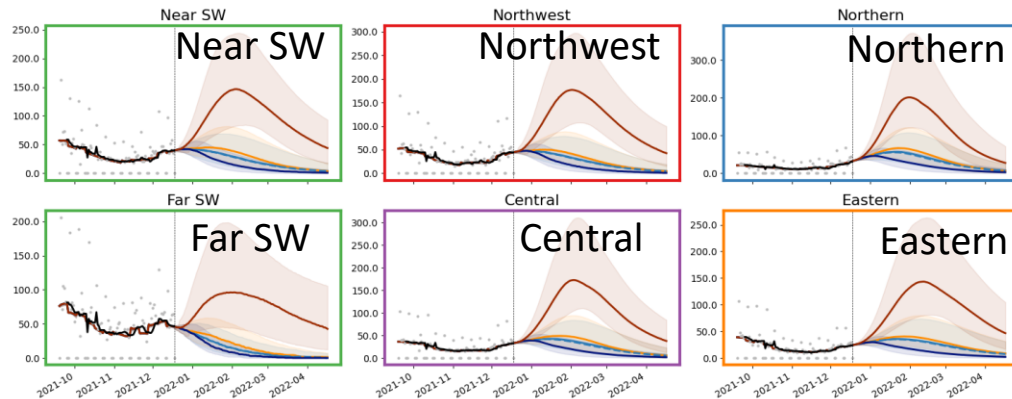


Daily confirmed cases)  
by rate (per 100K)  
District (grey with 7-day  
average in black) with  
simulation colored by  
scenario

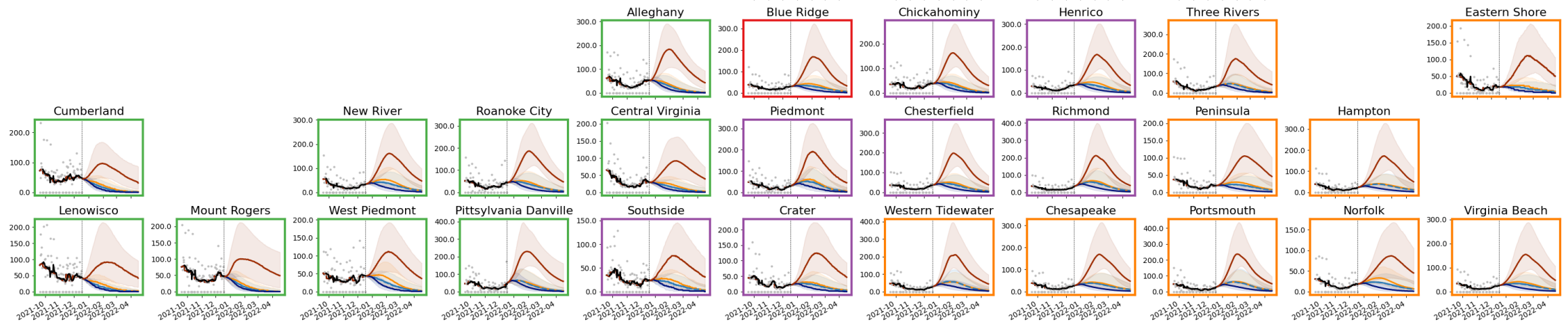


# Detailed Projections: All Scenarios - Closer Look

## Projections by Region



## Projections by District

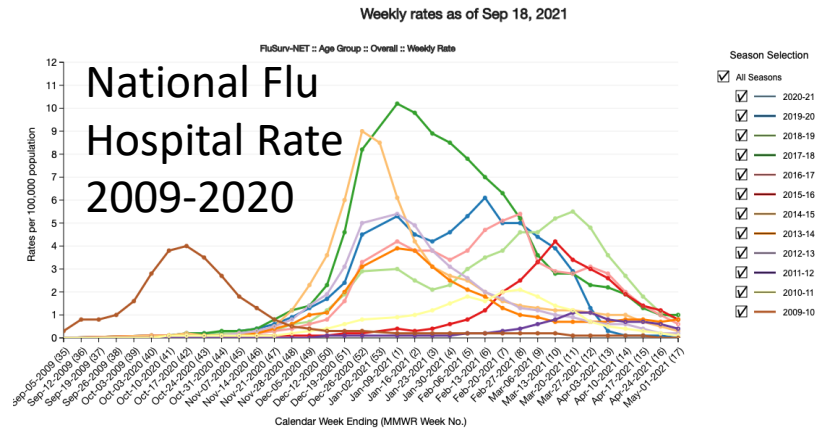


Daily confirmed cases by rate (per 100K) District (grey with 7-day average in black) with simulation colored by scenario

# Impact of Influenza based on Previous Intense Flu Seasons

## Augment COVID-19 daily hospitalizations with that of past Influenza seasons

- Include hybrid seasons that use timing of one season but are scaled by severity of another
- Due to limited historical data on Virginia flu hospitalizations currently using national rates applied to VA population



<https://gis.cdc.gov/GRASP/Fluview/FluHospRates.html>

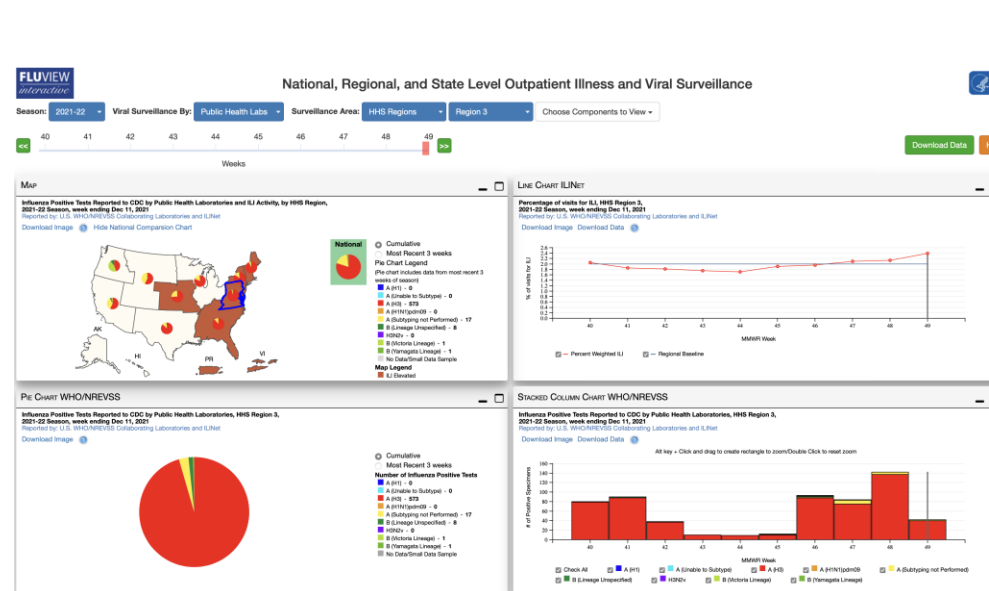
**2009-10** – Pandemic 2009 H1N1 season

**2017-18** – Timing and severity of 2017-18 season

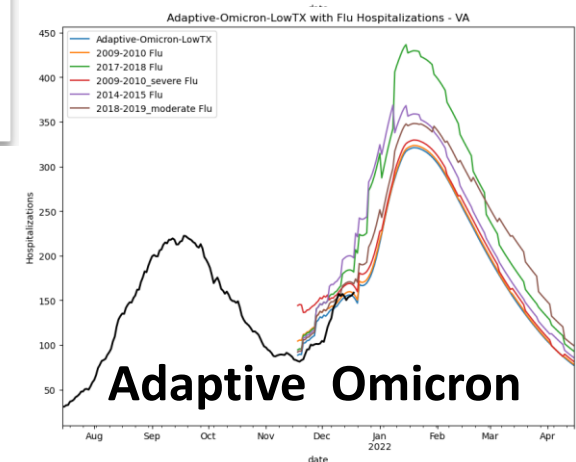
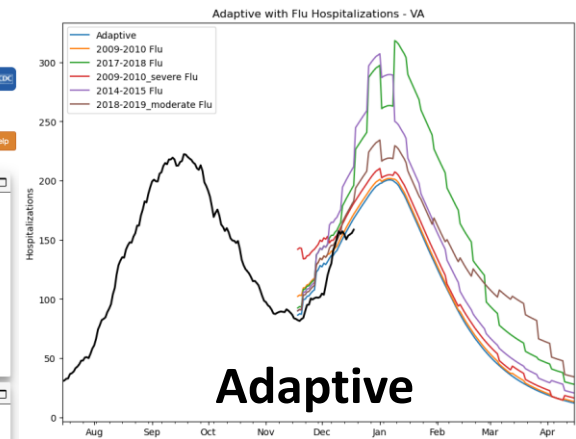
**2009-10\_severe** – Timing of 2009 pandemic (early) with the severity of the 2017-18 season

**2014-15** – Timing and severity of 2014-15 season

**2018-19\_moderate** – Timing of 2018-19 (late) season with severity of 2014-15



Influenza A activity up in our region  
Labs show high levels of H3 this season  
(Influenza A H3N2 is more severe)



# Key Takeaways

Projecting future cases precisely is impossible and unnecessary.

Even without perfect projections, we can confidently draw conclusions:

- **Case rates grew after holiday break but growth has slowed slightly, keeping case rates high as the anticipated arrival of Omicron may fuel more rapid growth in the near term**
- VA 7-day mean daily case rate up to 30/100K from 26/100K; US is up to 36/100K (from 35/100K)
- Projections show a continued rise of cases which becomes more extreme under Omicron and FallWinter scenarios that anticipate likely drivers of future transmission
- Recent updates:
  - Overhauled model structure further refined to better capture different tiers of immunity and the immune evasion of the Omicron variant
  - Analysis of the effects of increasing 3<sup>rd</sup> dose coverage

The situation continues to change. Models continue to be updated regularly.

# Additional Analyses

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# New Infections by Vaccine Status

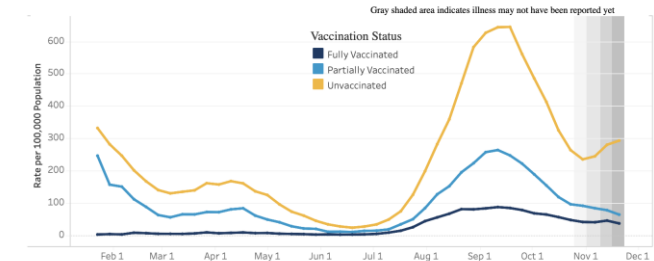
## Infections among Naïve, Vaccinated, and Partial Vaccinated

- VDH data for cases with known vaccination status show ~20-30% of current infections come from tiers with some vaccine induced immunity (~20% full and 5-10% partial)
- Model estimates of all infections are similar however, under the Omicron scenario the vast majority of future cases will be from those with prior immunity (due immune evasion)
- Infections in those with prior immunity are likely to be less severe (less hospitalizations and deaths)

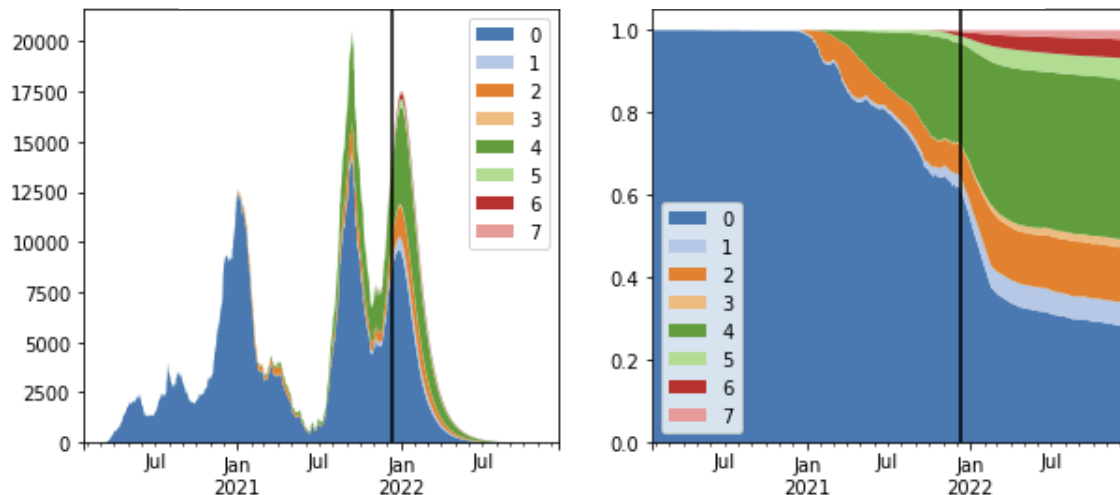
**COVID-19 in Virginia:  
Case Rates by Vaccination Status**  
Dashboard Updated: 11/26/2021 Data through: 11/20/2021  
Updated Weekly on Fridays

| Rates by Vaccination Status                             |                              |                     |
|---|------------------------------|---------------------|
| Select Metric   | Week Ending Date             |                     |
| Infections  | (All)                        |                     |
| Rate of Infections per 100,000*, 1/17/2021 - 11/20/2021 |                              |                     |
| Fully Vaccinated** People                               | Partially Vaccinated† People | Unvaccinated People |
| 1,098   | 2,227                        | 4,997               |

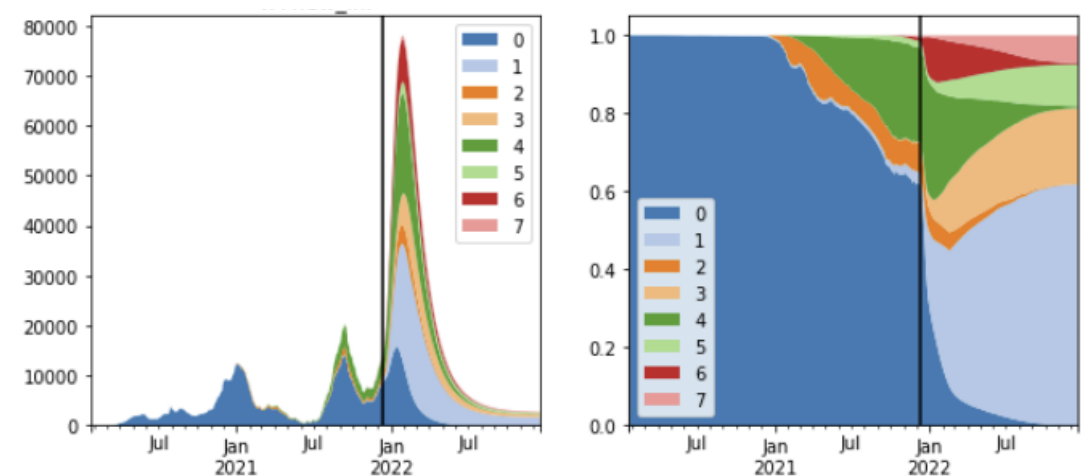
Between 1/17/2021 and 11/20/2021, unvaccinated people developed COVID-19 at a rate **4.6 times** that of fully vaccinated people and **2.2 times** that of partially vaccinated people.‡



Adaptive - New Infections by Immune Tier

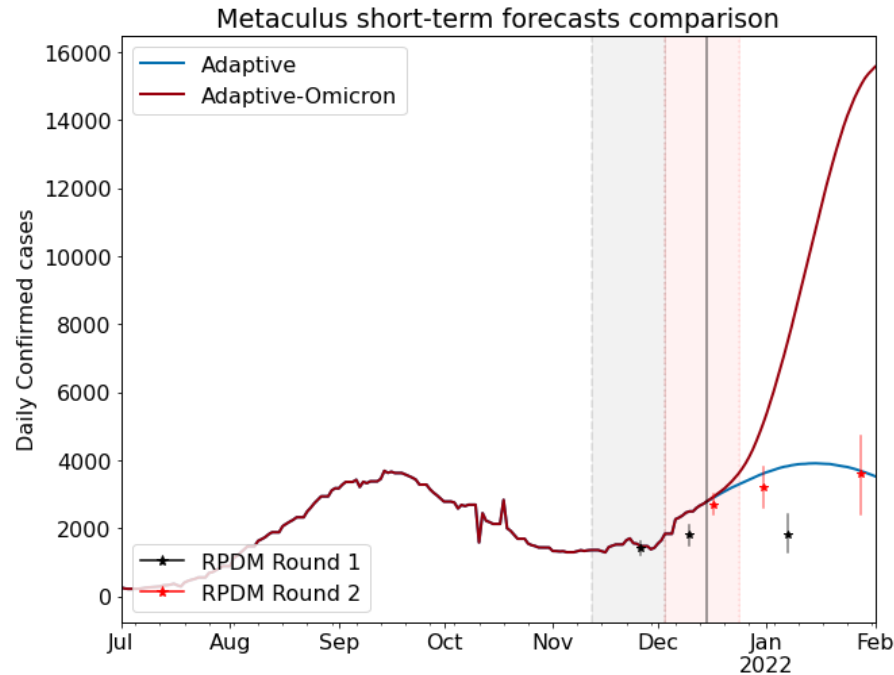


Adaptive-Omicron New Infections by Immune Tier



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# Metaculus - Short Term & Omicron Forecasts

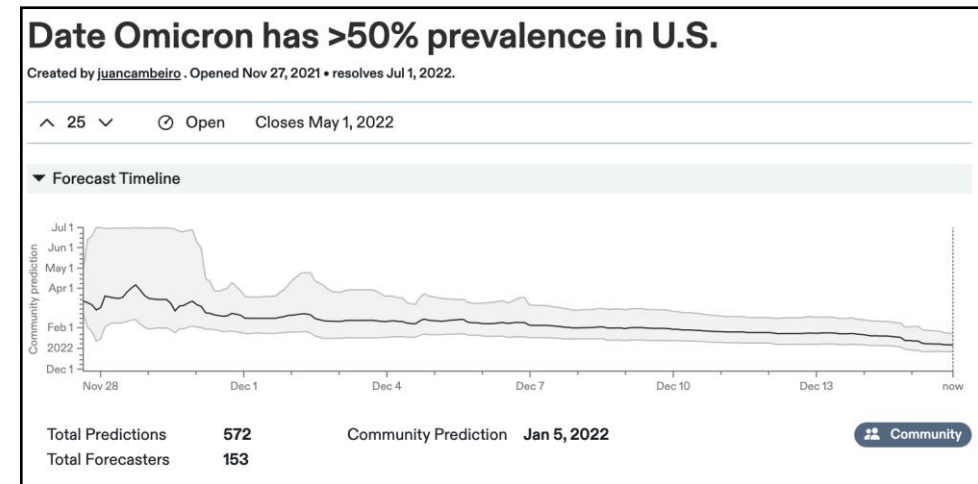
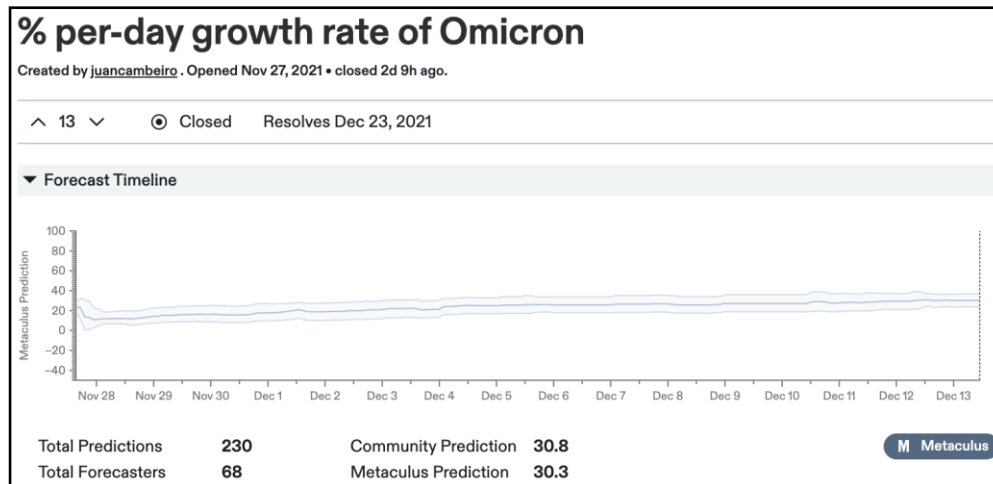


## Short-term<sup>1</sup>

- Round 1 forecasts underpredicted current surge
- Round 2 forecasts are more similar to current Adaptive, but much lower than Omicron scenario

## Omicron<sup>2</sup>

- Community estimate of growth rate is similar to current model input (**~31%**)
- Predicted midpoint for US (**January 5<sup>th</sup>, 2022**) is later (and trending downwards from early prediction)



# Overview of relevant on-going studies

Other projects coordinated with CDC and VDH:

- **Scenario Modeling Hub:** Consortium of academic teams coordinated via MIDAS / CDC to that provides regular national projections based on timely scenarios
- **Genomic Surveillance:** Analyses of genomic sequencing data, VA surveillance data, and collaboration with VA DCLS to identify sample sizes needed to detect and track outbreaks driven by introduction of new variants etc.
- **Mobility Data driven Mobile Vaccine Clinic Site Selection:** Collaboration with VDH state and local, Stanford, and SafeGraph to leverage anonymized cell data to help identify

# COVID-19 Scenario Modeling Hub

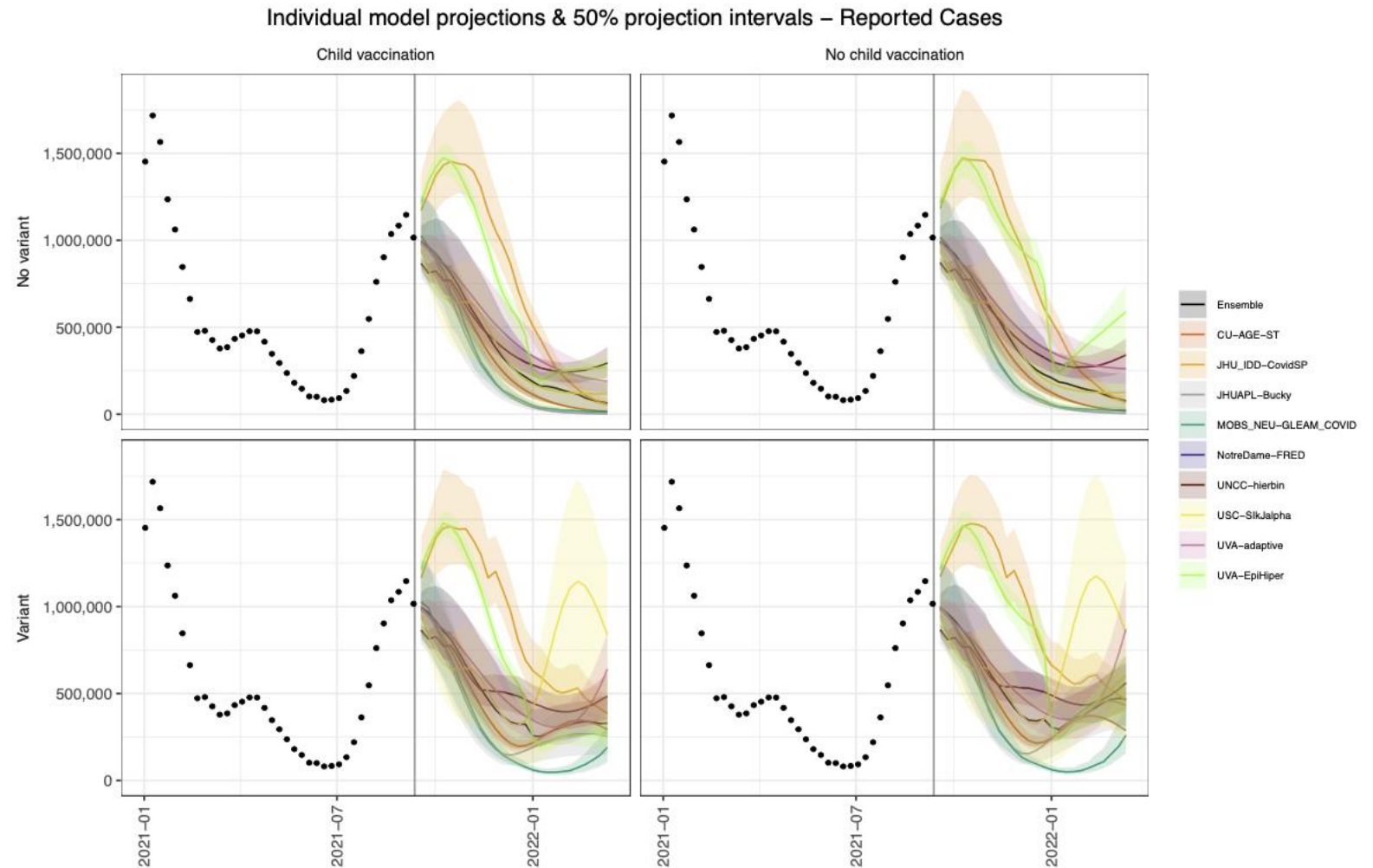
Collaboration of multiple academic teams to provide national and state-by-state level projections for 4 aligned scenarios that vary vaccine rates (high – low) and impact of the Delta variant (high and low)

- Round 9 released to assist in support of 5-11 vax consideration (ACIP meeting Sept 22-23)

- Rounds 4-8 now available

*Round 4 Results were published May 5<sup>th</sup>, 2021 in [MMWR](#)*

<https://covid19scenariomodelinghub.org/viz.html>





# COVID-19 Scenario Modeling Hub – Round 7

Round 7 scenarios explore the effects of a variant similar to Delta (B.1.617.2) against different backgrounds of vaccination. Includes some vax escape

## Vaccinations in 5-11 start in Nov

- Follows same rates as adolescents

## Emerging Variant Impact (5% prevalence on Nov 15)

- 50% boost as it eventually predominates

We consider a 2x2 scenario design, where childhood vaccination (5-11 years) is on the first axis, and a change in virus transmissibility is on the second axis. The second axis reflects a stress test, illustrating the potential impact of a new variant arising during the projection period:

|   |   |   |
|---|---|---|
|   | The same mix of variants circulate throughout the projection period. No change in virus transmissibility. | A more transmissible variant emerges, comprising 1% of circulating viruses on <b>Nov 15</b> . The new variant is <b>1.5X</b> as transmissible as viruses circulating at the beginning of the projection period. |
| Vaccination among 5-11yrs is approved and immunization begins on Nov 1. Each state's uptake rate reflects the percent coverage increases observed for 12-17-year-olds since distribution began on May 13. | A   | C   |
| No vaccination for children under 12  | B   | D   |

<https://covid19scenariomodelinghub.org/viz.html>

# Preliminary Analysis of Impact of Waning and 3<sup>rd</sup> doses

## Study to assess impact of waning rate and 3<sup>rd</sup> dose coverage levels

**Waning rate:** Duration population remains in an immune state (Vax or Recovered) until becoming susceptible

- Pessimistic: Mean duration 6 months
- Optimistic: Mean duration 1 year

**3<sup>rd</sup> Dose Coverage:** Proportion of Fully Vaccinated that receive a 3<sup>rd</sup> dose and return to full protection

- High: 70% coverage
- Low: 40% coverage

| Scenario           | Waning Rate | 3 <sup>rd</sup> Dose Coverage |
|--------------------|-------------|-------------------------------|
| A: optWan_highBoo  | 1 year      | 70%                           |
| B: optWan_lowBoo   | 1 year      | 40%                           |
| C: pessWan_highBoo | 6 months    | 70%                           |
| D: pessWan_lowBoo  | 6 months    | 40%                           |

### Partial Protection for:

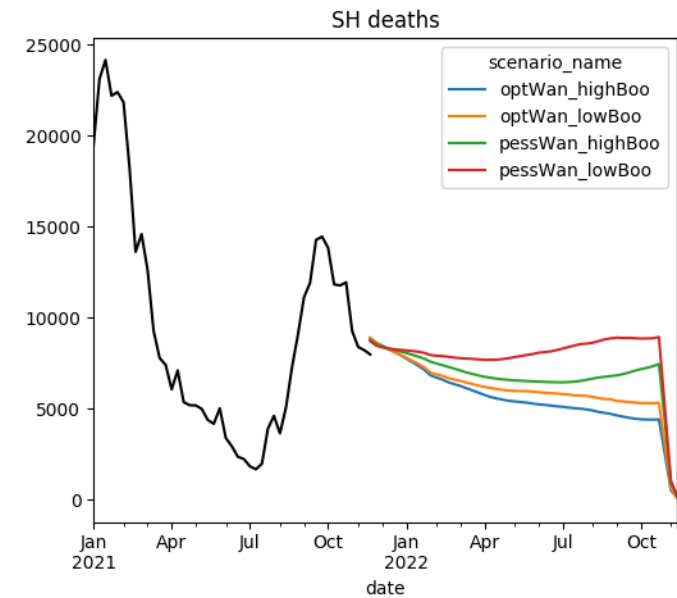
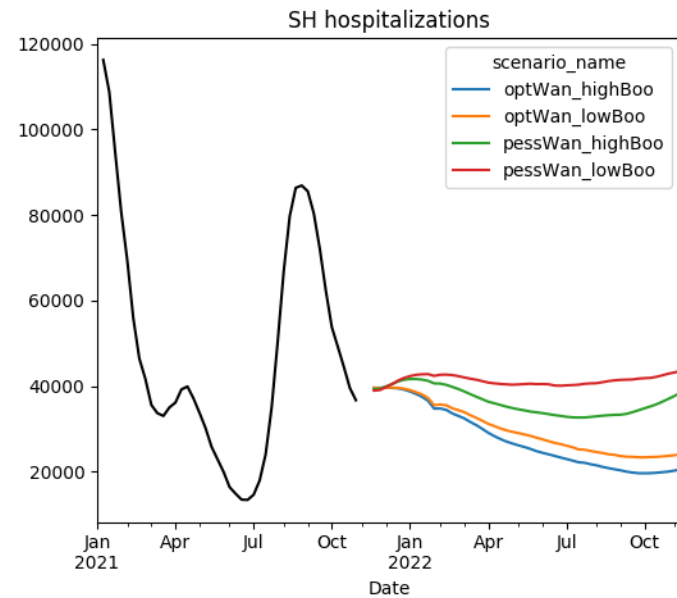
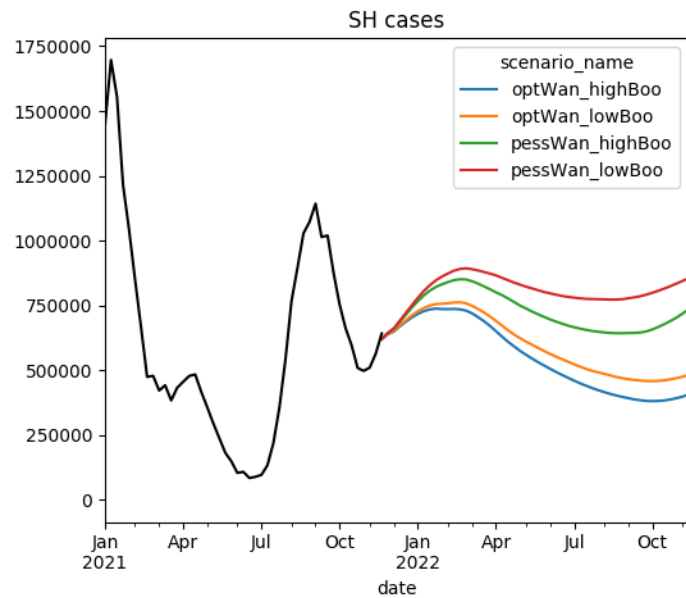
#### Optimistic Waning

| Protection against | Less than 65 | 65 + |
|--------------------|--------------|------|
| Infection          | 60%          | 40%  |
| Hospitalization    | 90%          | 80%  |
| Death              | 95%          | 90%  |

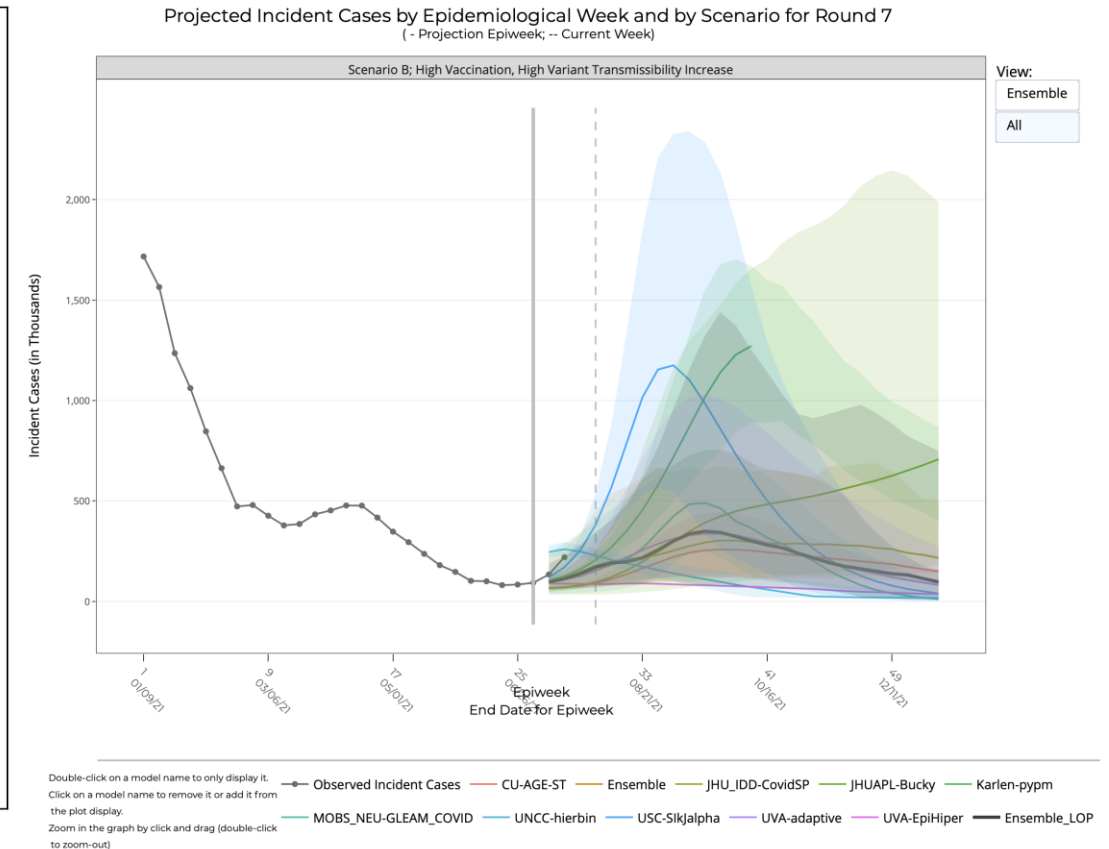
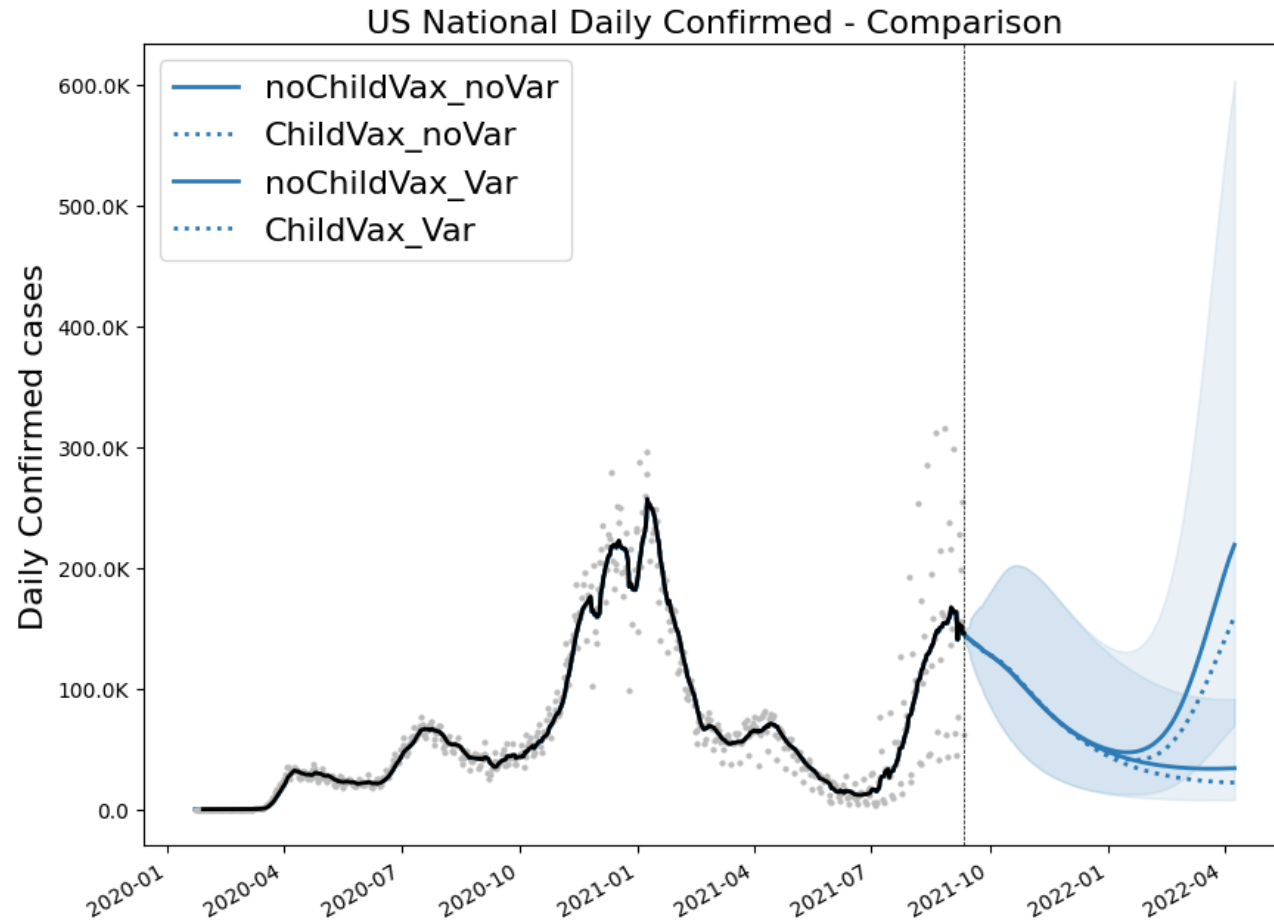
#### Pessimistic Waning

| Protection against | Less than 65 | 65 + |
|--------------------|--------------|------|
| Infection          | 50%          | 30%  |
| Hospitalization    | 80%          | 70%  |
| Death              | 90%          | 85%  |

# Preliminary Analysis of Impact of Waning and Boosters



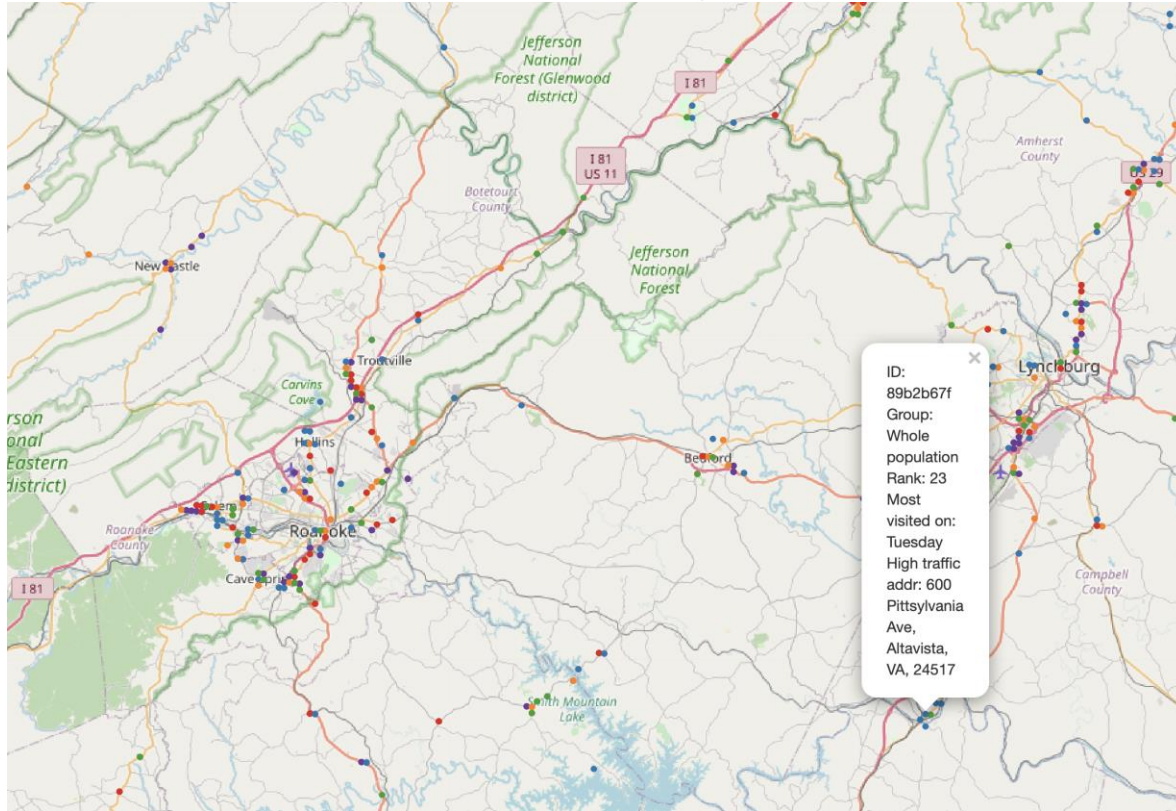
# Modeling Hub – Round 9 Prelim Results





# Data Recommended Mobile Vax Clinic Sites

## Detailed and Timely Locations



## Data Delivered and Disseminated to Locals

Provides a list of areas most visited by a given demographic group based on SafeGraph mobility data that links visits to specific sites and the home Census Block Group of the anonymized visitors

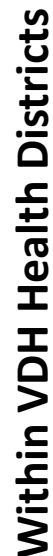
**Demographic Groups:** Black, Lantinx, Young Adults (20-40), Unvaccinated, and Whole Population

**Data Included:** Rank, Weight, most visited Day of Week, Highly Visited Address, and Lat-Long of area

**Goal:** Provide frequently visited locations based on populations and vaccination levels one desires to reach

**Example:** List of location in the Southside frequented by 20-40 year olds

## Overlap of locations between groups



- ## Different groups visit different areas
- Least overlap between Black and Latinx
  - Overlap in ages highest, but drops with large gap
  - Districts have different overlap patterns

# References

Venkatramanan, S., et al. "Optimizing spatial allocation of seasonal influenza vaccine under temporal constraints." *PLoS Computational Biology* 15.9 (2019): e1007111.

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NSSAC. PatchSim: Code for simulating the metapopulation SEIR model. <https://github.com/NSSAC/PatchSim>

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Biocomplexity Institute. COVID-19 Surveillance Dashboard. <https://nssac.bii.virginia.edu/covid-19/dashboard/>

Google. COVID-19 community mobility reports. <https://www.google.com/covid19/mobility/>

Biocomplexity page for data and other resources related to COVID-19: <https://covid19.biocomplexity.virginia.edu/>

# Questions?

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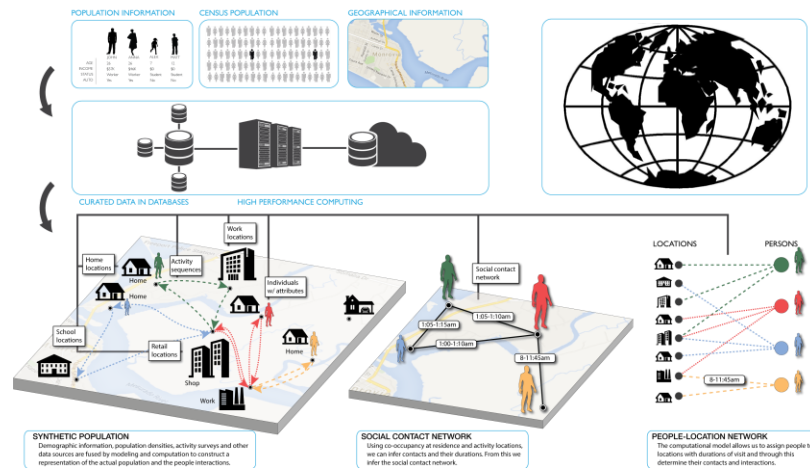


# Supplemental Slides

# Agent-based Model (ABM )

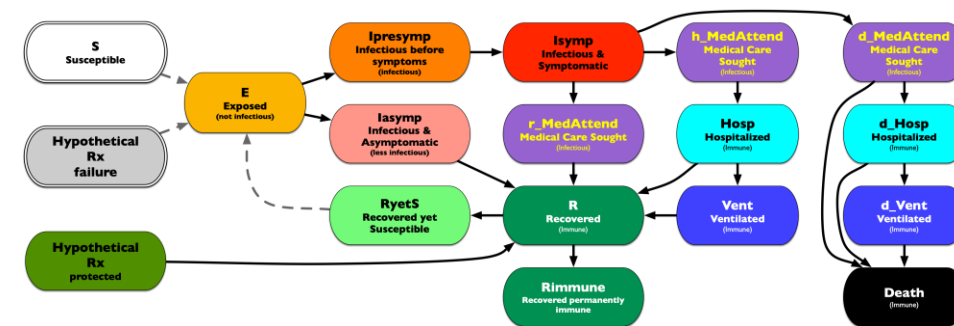
## EpiHiper: Distributed network-based stochastic disease transmission simulations

- Assess the impact on transmission under different conditions
- Assess the impacts of contact tracing



### Synthetic Population

- Census derived age and household structure
- Time-Use survey driven activities at appropriate locations



### Detailed Disease Course of COVID-19

- Literature based probabilities of outcomes with appropriate delays
- Varying levels of infectiousness
- Hypothetical treatments for future developments